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(54) Title: HEMATOPHAGOUS INSECT CALRETICULIN NUCLEIC ACID MOLECULES, PROTEINS AND USES THEREOF

(57) Abstract

The present invention is directed to novel insect calreticulin nucleic acid molecules and proteins. The present invention includes hematophagous insect nucleic acid molecules that hybridize under stringent hybridization conditions to a flea calreticulin gene as well as hematophagous insect calreticulin proteins encoded by such nucleic acid molecules. The present invention also includes therapeutic compositions capable of reducing insect infestation or allergic dermatitis, as well as methods to treat animals with such therapeutic compositions. Examples of such therapeutic compositions include hematophagous insect calreticulin proteins, nucleic acid molecules encoding such proteins, anti-calreticulin antibodies and calreticulin inhibitors.

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HEMATOPHAGOUS INSECT CALRETICULIN NUCLEIC ACID MOLECULES, PROTEINS AND USES THEREOF

FIELD OF THE INVENTION

The present invention relates to novel hematophagous insect calreticulin nucleic acid molecules, proteins encoded by such nucleic acid molecules, antibodies to such proteins, inhibitors of such proteins, and uses of therapeutic compositions comprising such nucleic acid molecules, proteins, antibodies and/or inhibitors to protect an animal from insect infestation, both on and in the environment surrounding the animal.

BACKGROUND OF THE INVENTION

Hematophagous (i.e., bloodsucking) insect infestation, and in particular flea infestation, of animals, is a health and economic concern because insects are known to cause and/or transmit a variety of diseases. Insects directly cause a variety of diseases, including allergies, and also carry a variety of infectious agents including, but not limited to, endoparasites (e.g., nematodes, cestodes, trematodes and protozoa), bacteria and viruses. In particular, the bites of insects are a problem for animals maintained as pets because the infestation becomes a source of annoyance not only for the pet but also for the pet owner who may find his or her home generally contaminated with insects. As such, insects are a problem not only when they are on an animal but also when they are in the general environment of the animal.

Bites from fleas are a particular problem because they not only can lead to disease transmission but also can

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cause a hypersensitive response in animals which is manifested in a disease called flea allergic (or allergy) dermatitis (FAD). A hypersensitive response in animals typically results in localized tissue inflammation and damage, causing substantial discomfort to the animal.

The medical and veterinary importance of insect infestation has prompted the development of reagents capable of controlling insect infestation. encountered methods to control insect infestation are generally focussed on use of insecticides in formulations such as sprays, shampoos, dusts, dips, or foams, or in pet collars. While some of these products are efficacious, most, at best, offer protection of a very limited duration. Furthermore, many of the methods are often not successful in reducing insect populations on the pet for one or more of the following reasons: (1) failure of owner compliance (frequent administration is required); (2) behavioral or physiological intolerance of the pet to the pesticide product or means of administration; and (3) the emergence of insect populations resistant to the prescribed dose of pesticide.

alternative method for controlling insect infestation is the use of insect vaccines administered to animals prior to or during infestation. However, despite considerable interest in developing anti-insect reagents, no insect presently exists of which the inventors are aware.

Calreticulins are calcium-binding proteins that have

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been found in the endoplasmic reticulum of a number of mammals (e.g., humans, rabbits, rats, mice), endoparasites (e.g., Schistosoma mansoni and Onchocerca volvulus) and even in the barley plants. In contrast, the inventors are not aware of any reports in the literature identifying calreticulin as a component of insect saliva, and in fact, the only saliva in which calreticulin has been found is in tick saliva (Jaworski, Ph.D. Dissertation, Ohio State University, Volume 52/08-B of Dissertation Abstracts International, 1991 and Jaworski et al., Biochem. Cell Biol., 71:11-12, 1993). That calreticulin is a component of insect saliva is unexpected even in view of the report of calreticulin in tick saliva not only because insects are phylogenetically different from ticks and other arachnids, but also because insects arose as a monophyletic order within the Class Insecta approximately 150 million years Furthermore, insects in general, and fleas in particular, feed from their hosts in a manner quite different from ticks. For example, fleas feed frequently and quickly, usually over minutes, while ticks spend several days preparing a bite wound before beginning to ingest blood and will feed uninterrupted for 5-7 days. Thus, the biological activity of components of flea saliva would be expected to differ from components of tick saliva. Indeed, studies by Jaworski, 1991, ibid., indicate that administration of calreticulin to a tick host causes necrosis and hemorrhaging at the site of tick feeding.

In summary, there remains a need to develop a reagent

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and a method to protect animals from insect infestation and diseases caused by such infestation.

SUMMARY OF THE INVENTION

The present invention relates to, in one embodiment, 5 an isolated hematophagous insect calreticulin protein encoded by a nucleic acid molecule that hybridizes under stringent hybridization conditions to a flea calreticulin gene. A preferred flea calreticulin gene comprises nucleic acid sequences SEQ ID NO:1, SEQ ID NO:11, SEQ ID NO:13, SEQ ID NO:14, SEQ ID NO:15, and SEQ ID NO:16; as such, a 10 preferred flea calreticulin gene encodes a protein comprising amino acid sequences SEQ ID NO:2 and SEQ ID NO:12. Preferably, the insect calreticulin is derived from insects including fleas, midges, mosquitos, sand flies, black flies, horse flies, horn flies, deer flies, tsetse 15 flies, buffalo flies, blow flies, stable flies, myiasiscausing flies, biting gnats, lice and true bugs, more preferably fleas of a genus including Ctenocephalides, Cyopsyllus, Diamanus, Echidnophaga, Nosopsyllus, Pulex, Tunga, Oropsylla, Orchopeus or Xenopsylla, and even more 20 preferably a species including Ctenocephalides felis, Ctenocephalides canis, Pulex irritans, Oropsylla (Thrassis) bacchi, Oropsylla (Diamanus) montana, Orchopeus howardi, Xenopsylla cheopis or Pulex simulans.

Another embodiment of the present invention includes an isolated hematophagous insect nucleic acid molecule that hybridizes under stringent hybridization conditions to a flea calreticulin gene. In particular, the nucleic acid

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molecule comprises a nucleic acid sequence having at least about 80% identity with nucleic acid sequence SEQ ID NO:1, SEQ ID NO:11, SEQ ID NO:13, SEQ ID NO:14, SEQ ID NO:15, and/or SEQ ID NO:16. Also included in the present invention are recombinant molecules and recombinant cells having a nucleic acid molecule of the present invention.

Yet another embodiment of the present invention includes a therapeutic composition for protecting an animal from hematophagous insect infestation, the composition comprising compounds including an isolated hematophagous insect calreticulin protein or a mimetope thereof, an isolated nucleic acid molecule capable of hybridizing under stringent hybridization conditions with a flea calreticulin gene, an isolated anti-hematophagous insect calreticulin antibody or a mimetope thereof, and/or a calreticulin inhibitory compound. The composition, when administered to an animal, is able to reduce hematophagous insect burden on the animal and in the environment of the animal. Preferred animals to protect with a therapeutic composition of the present invention include mammals and birds, with cats, dogs, sheep, cows, pigs, horses and goats being more preferred. The present invention also relates to a method to protect an animal from insect infestation, comprising treating an animal with a therapeutic composition of the present invention.

One aspect of the present invention includes a method to produce a hematophagous insect calreticulin protein comprising culturing in an effective medium a recombinant

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cell transformed with a nucleic acid molecule encoding the protein to produce the protein.

Another aspect of the present invention includes a method to identify a compound capable of inhibiting hematophagous insect calreticulin activity, said method comprising: (a) contacting an isolated hematophagous insect calreticulin protein with a putative inhibitory compound under conditions in which, in the absence of the compound, the protein has calreticulin activity; and (b) determining if the putative inhibitory compound inhibits said calreticulin activity. The present invention also includes test kits to identify inhibitory compounds and inhibitory compounds identified using a test kit and/or method of the present invention.

DETAILED DESCRIPTION OF THE INVENTION

The present invention includes hematophagous insect calreticulin nucleic acid molecules, proteins encoded by such nucleic acid molecules, antibodies raised against such proteins, inhibitors of such proteins and the use of therapeutic compositions comprising such nucleic acid molecules, proteins, antibodies and/or inhibitors (i.e., inhibitory compounds) to protect an animal from insect infestation and insect allergic dermatitis. The present invention is particularly advantageous in that, since calreticulin can be found in the saliva of insects, therapeutic compositions of the present invention can be used to alter the ability of hematophagous insects to acquire and digest blood meals from an animal treated with

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the composition. A primary function of calreticulin is to regulate calcium concentrations, such as concentrations that regulate calcium-dependent blood coagulation. Without being bound by theory, it is believed that calreticulin is produced and secreted by the salivary gland of an insect as the insect feeds from a host and is washed into the gut with the blood meal. During blood meal acquisition, therefore, the secreted calreticulin is believed to influence blood coagulation and blood vessel dilation at the feeding site by, for example, interfering with platelet aggregation and/or altering nitric oxide synthetase production. During blood meal digestion in the gut, calreticulin can also reduce blood clotting by, for example modifying the activity of components (e.g., blood factors) of the blood coagulation pathways.

One embodiment of the present invention is an isolated hematophagous insect calreticulin protein encoded by a nucleic acid molecule that hybridizes under stringent hybridization conditions to a flea calreticulin gene. According to the present invention, hematophagous insects are external living insects that attach and feed through the skin of a host animal. Hematophagous insects can live on a host animal or attach temporarily to an animal in order to feed.

As used herein, a flea calreticulin gene includes all nucleic acid sequences related to a natural flea calreticulin gene such as regulatory regions that control production of a flea calreticulin protein encoded by that

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but not limited to, transcription, gene (such as, translation or post-translation control regions) as well as the coding region itself. A flea calreticulin gene can be an isolated natural flea calreticulin gene or a homologue thereof. A flea calreticulin gene can be included in a nucleic acid molecule encoding a hybrid protein, a fusion protein, a multivalent protein or a truncation fragment. As used herein, stringent hybridization conditions refer to standard hybridization conditions under which nucleic acid molecules are used to identify similar nucleic acid Additional definition and examples of such molecules. conditions are provided herein.

A preferred flea calreticulin gene of the present

invention encodes a protein that includes, but is not 15 limited to, the amino acid sequence represented herein as SEQ ID NO:2. Also preferred is a flea calreticulin gene encoding a protein having the amino acid sequence SEQ ID NO:12, which apparently represents the full-length protein. A more preferred flea calreticulin gene of the present invention includes, but is not limited to, the nucleic acid 20 sequence referred to herein as SEQ ID NO:1 and/or the following nucleic acid sequences: SEQ ID NO:11, SEQ ID NO:13, SEQ ID NO:14, SEQ ID NO:15, and/or SEQ ID NO:16. The determination of SEQ ID NO:1 and SEQ ID NO:2, as well as other sequences presented herein, is described in the 25 Examples. The included SEQ ID NO's represent nucleic acid and amino acid sequences deduced according to methods disclosed in the Examples. It should be not d that since

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nucleic acid and amino acid sequencing technology is not entirely error-free, the fore-going SEQ ID NO's, at best, represent an apparent nucleic acid sequence of the isolated nucleic acid molecule used to obtain SEQ ID NO:1 or an amino acid sequence deduced from SEQ ID NO:1 (i.e., SEQ ID NO:2), or other sequences presented herein.

Suitable flea calreticulin genes of the present invention encode calreticulin proteins that can be isolated from flea saliva. Flea saliva refers to the material released from the mouth of a flea when the flea attempts to feed in response to a temperature differential between the skin temperature of a host animal and the temperature of the air surrounding the host, or in response to an artificially created temperature differential such as exists in an artificial flea feeding apparatus.

According to the present invention, an isolated hematophagous insect calreticulin protein is a hematophagous insect calreticulin protein that has been removed from its natural milieu. An isolated hematophagous insect calreticulin protein can, for example, be obtained from its natural source, be produced using recombinant DNA technology, or be synthesized chemically. As used herein, an isolated hematophagous insect calreticulin protein can be a full-length protein or a homologue thereof.

In one embodiment, an isolated hematophagous insect calreticulin protein of the present invention is a protein isolated from the saliva of an insect. As noted above, such an "isolated" protein can be a natural protein, a

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synthetic protein, or a recombinant protein.

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A preferred isolated hematophagous insect calreticulin protein of the present invention is a protein that, when administered to an animal, is capable of protecting that animal from insect infestation and/or allergic dermatitis. As used herein, the phrase "to protect an animal from insect infestation" refers to reducing the potential for insect population expansion on and around an animal (i.e., reducing the insect burden). At any given time, a certain percentage of an insect population can be on a host animal whereas the remainder can be in the environment surrounding the animal (i.e., in the environment of the animal). host animal, as used herein, is an animal from which insects can feed. The environment can be of any size such that insects in the environment are able to attach onto and detach from a host animal. Preferably, the insect population size is decreased, optimally to an extent that the animal is no longer bothered by such insects. As such, it is desirable not only to reduce the insect burden on an animal per se, but also to reduce the insect burden in the environment surrounding the animal.

A preferred hematophagous insect calreticulin protein of the present invention can protect an animal from insect infestation by reducing calreticulin activity at the feeding site or in an insect (e.g., in the midgut) feeding from an animal administered such a protein. Such a reduction in calreticulin activity can be the result of the ability of the protein to reduce calcium binding activity

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and/or other activities of the insect calreticulin by, for example, eliciting an immune response against calreticulin in the insect feeding from the animal. As such, a preferred hematophagous insect calreticulin protein of the present invention can include at least one epitope capable of eliciting production of an antibody capable of binding to an insect calreticulin protein. It is to be noted that other calreticulin activities include but are not limited to, (a) ability to bind to other factors such as magnesium, cellular integrins, extracellular matrix proteins, DNA domains of nuclear hormone receptors, and/or resident endoplasmic reticulum proteins; (b) ability to affect protein translocation from an endoplasmic reticulum; and/or (c) ability to act as an agonist or antagonist of other calcium binding proteins.

Administration of an isolated hematophagous insect calreticulin protein of the present invention can (a) alter the ability of a hematophagous insect to feed from an animal (e.g., by altering the volume per feeding or altering the time and frequency of feeding) as a result of interfering with platelet aggregation or vasodilation; (b) interfere with blood coagulation at the feeding site or in the midgut of an insect that feeds from an animal; (c) interfere with the activation of blood complement ingested from an animal in the midgut of an insect; (d) reduce the number of eggs produced and/or their subsequent viability; and/or (e) modulate the immune system of an animal by regulating, for example, the activation of cells involved

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in an immune response, chemotaxis, lymphocyte responsiveness, and cytokine production. As used herein, the ability of a hematophagous insect calreticulin protein to alter insect feeding behavior refers to the protein's ability to alter blood meal acquisition and/or blood meal digestion.

In one embodiment, a hematophagous insect calreticulin protein of the present invention includes a protein that, when administered to an animal, is capable of substantially desensitizing the animal to allergic dermatitis. Preferred types of allergic dermatitis to desensitize an animal against include flea allergic dermatitis, Culicoides allergic dermatitis and mosquito allergic dermatitis, with flea allergic dermatitis being more preferred. herein, "desensitizing an animal" refers to treating an animal in such a manner that the animal does not exhibit a hypersensitive response to an allergen. Hypersensitivity refers to a state of altered reactivity in which an animal, having been previously exposed to a compound, exhibits an allergic compound upon subsequent response to the exposures. The term "allergen" primarily refers to foreign compounds capable of causing an allergic response. term can be used interchangeably with the term "antigen," especially with respect to a foreign compound capable of symptoms immediate and/or delayed inducing of hypersensitivity. Preferably, an animal is substantially desensitized to allergic dermatitis when the animal is blocked from having a hypersensitive response (e.g., edema,

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erythema, itching, inflammation) to the bite of an insect.

The ability of a hematophagous insect calreticulin protein to protect an animal from infestation or allergic dermatitis, can be tested using techniques known to those skilled in the art. Methods to measure insect infestation include determining insect viability, fecundity of female insects, reproductive capacity of male insects, viability of insect eggs, blood feeding behavior, viability of insect larvae, and/or development and maturation of larvae into adults. Techniques to test allergic dermatitis include standard allergen skin tests (e.g., IDST), immunoabsorbent assays (e.g., ELISA, Western blot, radioimmunoprecipitation assay), and passive cutaneous anaphylaxis.

heretofore hematophagous As stated, а insect 15 calreticulin protein of the present invention can be a full-length natural protein or a homologue thereof. used herein, a homologue can be a hematophagous insect calreticulin protein in which amino acids have been deleted (e.g., a truncated version of the protein, such as a 20 peptide), inserted, inverted, substituted and/or (e.g., by glycosylation, phosphorylation, acetylation, myristoylation, prenylation, palmitoylation, amidation and/or addition of glycosylphosphatidyl inositol). A homologue of a hematophagous 25 calreticulin protein is a protein having an amino acid sequence that is sufficiently similar to a natural hematophagous insect calreticulin protein amino sequence that a nucleic acid sequence encoding the

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homologue is capable of hybridizing under stringent conditions to (i.e., with) a nucleic acid sequence encoding the natural hematophagous insect calreticulin protein amino acid sequence. As used herein, stringent hybridization conditions refer to standard hybridization conditions under which nucleic acid molecules, including oligonucleotides, are used to identify similar nucleic acid molecules. Such standard conditions are disclosed, for example, in Sambrook et al., Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Labs Press, 1989. The minimal size of a protein homologue of the present invention is a size sufficient to be encoded by a nucleic acid molecule capable of forming a stable hybrid with the complementary sequence of a nucleic acid molecule encoding the corresponding natural protein. As such, the size of the nucleic acid molecule encoding such a protein homologue is dependent on nucleic acid composition and percent homology between the nucleic acid molecule and complementary sequence as well as upon hybridization conditions per se (e.g., temperature, salt concentration, and formamide concentration). The minimal size of such nucleic acid molecules is typically at least about 12 to about 15 nucleotides in length if the nucleic acid molecules are GC-rich and at least about 15 to about 17 bases in length if they are AT-rich. As such, the minimal size of a nucleic acid molecule used to encode a hematophagous insect calreticulin protein homologue of the present invention is from about 12 to about 18 nucleotides There is no limit, other than a practical in length.

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limit, on the maximal size of such a nucleic acid molecule in that the nucleic acid molecule can include a portion of a gene, an entire gene, or multiple genes, or portions thereof. Similarly, the minimal size of a hematophagous insect calreticulin protein homologue of the present invention is from about 4 to about 6 amino acids in length, but can be of any longer length depending on whether a full-length, multivalent (i.e., protein having more than one domain each of which has a protective function), fusion, hybrid or functional portions of such proteins are desired.

Hematophagous insect calreticulin protein homologues can be the result of allelic variation of a natural gene encoding a hematophagous insect calreticulin protein. A natural gene refers to the form of the gene found most often in nature. Hematophagous insect calreticulin protein homologues can be produced using techniques known in the art including, but not limited to, direct modifications to a gene encoding a protein using, for example, classic or recombinant DNA techniques to effect random or targeted mutagenesis.

Preferred hematophagous insect calreticulin protein homologues of the present invention are capable of protecting an animal from hematophagous insect infestation or allergic dermatitis resulting from the bites of insects in a similar manner as the natural hematophagous insect calreticulin protein counterpart. The ability of a hematophagous insect calreticulin protein homologue to

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protect an animal from hematophagous insect infestation or allergic dermatitis can be tested by the methods disclosed herein.

A preferred hematophagous insect calreticulin protein of the present invention comprises (i.e., includes, but is not limited to) an amino acid sequence that is at least about 85%, more preferably at least about 90%, and even more preferably at least about 95% identical to an amino acid sequence represented by SEQ ID NO:2 and/or to amino acid sequences spanning amino acids from about 10 through about 360 of SEQ ID NO:12. A more preferred hematophagous insect calreticulin protein of the present invention comprises an amino acid sequence encoded by a nucleic acid molecule that is at least about 80%, more preferably at least about 85%, even more preferably at least about 90%, and even more preferably at least about 95% identical to a nucleic acid sequence represented by SEQ ID NO:1, and/or to a nucleic acid sequence spanning about nucleotide 360 through about 860 of SEQ ID NO:11. For example, Meinkoth et al, 1984, Anal. Biochem 138, 267-284, provide formulae to calculate the appropriate hybridization and wash conditions to achieve hybridization at various degrees of nucleic acid identity as disclosed herein. al., ibid., is incorporated by reference herein in its entirety.

In a preferred embodiment, a hematophagous insect calreticulin protein of the present invention comprises an amino acid sequence represented by SEQ ID NO:2 and/or SEQ

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ID NO:12. particular, In a hematophagous calreticulin protein of the present invention comprises an amino acid sequence encoded by a nucleic acid sequence represented by SEQ ID NO:1, SEQ ID NO:11 and/or SEQ ID NO:14. A preferred embodiment includes flea calreticulin proteins PCtCal₈₅, PHIS-PCtCal₂₅₇, PCtCal₄₀₃ and PHIS-PCtCal₄₀₃, the production of which is described in the Examples.

As will be apparent to one of skill in the art, the present invention is intended to apply to all hematophagous insects. A preferred insect of the present invention from which to obtain hematophagous insect calreticulin proteins, and/or from which to identify proteins that can then be produced recombinantly or synthetically, include fleas; flies, such as midges (e.g., Culicoides), mosquitos, sand flies, black flies, horse flies, horn flies Haematobia irritans irritans), deer flies, tsetse flies, buffalo flies, blow flies, stable flies, myiasis-causing flies and biting gnats; lice and true bugs (e.g., Rhodinus and Cimex), such as bed bugs and kissing bugs, including 20 those carrying Chagas disease. A more preferred insect includes a flea, a myiasis-causing fly and a horn fly. A preferred flea of the present invention includes a flea being of the genus Ctenocephalides, Cyopsyllus, Diamanus, Echidnophaga, Nosopsyllus, Pulex, Tunga, Oropsylla, Orchopeus or Xenopsylla. A more preferred flea of the present invention includes a flea being of the species Ctenocephalides felis, Ctenocephalides canis, Pulex irritans, Oropsylla (Thrassis) bacchi, Oropsylla (Diamanus)

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montana, Orchopeus howardi, Xenopsylla cheopis or Pulex simulans.

Another embodiment of the present invention is an isolated hematophagous insect calreticulin nucleic acid molecule that hybridizes under stringent hybridization conditions to a flea calreticulin gene. In accordance with the present invention, an isolated nucleic acid molecule is a nucleic acid molecule that has been removed from its natural milieu (i.e., that has been subject to human manipulation). As such, "isolated" does not reflect the extent to which the nucleic acid molecule has been purified. An isolated nucleic acid molecule can include DNA, RNA, or derivatives of either DNA or RNA.

An isolated nucleic acid molecule of the present invention can be obtained from its natural source either as an entire (i.e., complete) gene or a portion thereof capable of forming a stable hybrid with that gene. As used herein, the phrase "at least a portion of" an entity refers to an amount of the entity that is at least sufficient to have the functional aspects of that entity. For example, at least a portion of a nucleic acid sequence, as used herein, is an amount of a nucleic acid sequence capable of forming a stable hybrid with the corresponding gene under stringent hybridization conditions. An isolated nucleic acid molecule of the present invention can also be produced using recombinant DNA technology (e.g., polymerase chain amplification, cloning) or reaction (PCR) Isolated hematophagous insect calreticulin synthesis.

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nucleic acid molecules include natural nucleic acid molecules and homologues thereof, including, but not limited to, natural allelic variants (the definition of which is known to those skilled in the art) and modified nucleic acid molecules in which nucleotides have been inserted, deleted, substituted, and/or inverted in such a manner that such modifications do not substantially interfere with the nucleic acid molecule's ability to encode a hematophagous insect calreticulin protein of the present invention or to form stable hybrids under stringent conditions with natural nucleic acid molecule isolates.

An isolated nucleic acid molecule of the present invention can include a nucleic acid sequence that encodes at least one hematophagous insect calreticulin protein of the present invention, examples of such proteins being disclosed herein. Although the phrase "nucleic acid molecule" primarily refers to the physical nucleic acid molecule and the phrase "nucleic acid sequence" primarily refers to the sequence of nucleotides on the nucleic acid molecule, the two phrases can be used interchangeably, especially with respect to a nucleic acid molecule, or a nucleic acid sequence, being capable of encoding a hematophagous insect calreticulin protein. As heretofore disclosed, a hematophagous insect calreticulin protein of the present invention include, but are not limited to, proteins having full-length hematophagous calreticulin protein coding regions and homologues thereof.

As heretofore disclosed, a hematophagous insect

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calreticulin gene includes all nucleic acid sequences related to a natural hematophagous insect calreticulin gene such as regulatory regions that control production of a hematophagous insect calreticulin protein encoded by that gene (such as, but not limited to, transcription, translation or post-translation control regions) as well as the coding region itself. A nucleic acid molecule of the present invention can be an isolated natural hematophagous insect calreticulin nucleic acid molecule or a homologue thereof. A nucleic acid molecule of the present invention can include one or more regulatory regions, full-length or partial coding regions, or combinations thereof. minimal size of a hematophagous insect calreticulin nucleic acid molecule of the present invention is the minimal size capable of forming a stable hybrid under stringent hybridization conditions with a corresponding natural gene.

A hematophagous insect calreticulin nucleic acid molecule homologue can be produced using a number of methods known to those skilled in the art (see, for example, Sambrook et al., ibid.). For example, nucleic acid molecules can be modified using a variety of limited to, techniques including, but not mutagenesis techniques and recombinant DNA techniques, such as site-directed mutagenesis, chemical treatment of a nucleic acid molecule to induce mutations, restriction enzyme cleavage of a nucleic acid fragment, ligation of nucleic acid fragments, polymerase chain reaction (PCR) amplification and/or mutagenesis of selected regions of a

nucleic acid sequence, synthesis of oligonucleotide mixtures and ligation of mixture groups to "build" a mixture of nucleic acid molecules and combinations thereof. Nucleic acid molecule homologues can be selected from a mixture of modified nucleic acids by screening for the function of the protein encoded by the nucleic acid (e.g., the ability of a homologue to protect an animal from insect infestation or from allergic dermatitis) and/or by hybridization with an isolated hematophagous insect calreticulin gene under stringent conditions.

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A preferred nucleic acid molecule of the present invention includes a nucleic acid molecule that encodes an amino acid sequence having at least about 85%, more preferably at least about 90%, and even more preferably at least about 95% identity to amino acid sequence represented as SEQ ID NO:2, or to amino acid sequences spanning amino acids from about 10 through about 360 of SEQ ID NO:12. A more preferred nucleic acid molecule comprises a nucleic acid sequence having at least about 80%, more preferably at least about 85%, even more preferably at least about 90%, and even more preferably at least about 95% identity to a nucleic acid sequence represented as SEQ ID NO:1 or to a nucleic acid sequence spanning about nucleotide 360 through about 860 of SEQ ID NO:11, or to the complements of those sequences.

A particularly preferred nucleic acid molecule of the present invention includes a nucleic acid sequence encoding an amino acid sequence represented herein as SEQ ID:2

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and/or SEQ ID NO:12; and/or a nucleic acid molecule that includes a nucleic acid sequence represented herein as SEQ ID NO:1, SEQ ID NO:11, SEQ ID NO:13, SEQ ID NO:14, SEQ ID NO:15, and/or SEQ ID NO:16. One preferred embodiment includes nucleic acid molecules nctcal₂₅₇, nctcal₈₅₀, nctcal₅₅₀, nctcal₅₅₆, nctcal₁₅₈₉, nctcal₁₂₀₉, nctcal₁₂₁₂, nctcal₆₆₅, nctcal₇₅₀, and nctcal₁₂₁₈.

Knowing a nucleic acid molecule of a hematophagous insect calreticulin protein of the present invention allows one skilled in the art to make copies of that nucleic acid molecule as well as to obtain a nucleic acid molecule including additional portions of hematophagous insect calreticulin protein-encoding genes (e.g., nucleic acid molecules that include a full-length coding region and/or transcription and/or translation control regions), and/or hematophagous insect calreticulin nucleic acid molecule homologues. Knowing a portion of an amino acid sequence of a hematophagous insect calreticulin protein of the present invention allows one skilled in the art to clone nucleic acid sequences encoding such a hematophagous insect Additional desired hematophagous calreticulin protein. insect calreticulin nucleic acid molecules can be obtained in a variety of ways including, but not limited to, screening appropriate expression libraries with antibodies which bind to a hematophagous insect calreticulin protein of the present invention; use of oligonucleotide probes of the present invention to screen appropriate libraries or DNA; and PCR amplification of appropriate libraries, or RNA

or DNA using oligonucleotide primers of the present invention (genomic and/or cDNA libraries can be used). To isolate a hematophagous insect calreticulin nucleic acid molecule, preferred cDNA libraries include cDNA libraries made from insect salivary glands. Techniques to clone and amplify genes are disclosed, for example, in Sambrook et al., ibid. The Examples section includes examples of the isolation of a cDNA sequence encoding a hematophagous insect calreticulin protein of the present invention.

10 The present invention also includes nucleic acid molecules that are oligonucleotides capable of hybridizing, under stringent conditions, with complementary regions of other, preferably longer, nucleic acid molecules of the present invention such as a flea calreticulin gene. preferred oligonucleotide is capable of hybridizing, under 15 stringent conditions, with a nucleic acid molecule that encodes a protein comprising SEQ ID NO:2 and/or SEQ ID NO:12, complements thereof. or Α more preferred oligonucleotide is capable of hybridizing, under stringent 20 conditions, to a nucleic acid molecule comprising SEQ ID NO:1, SEQ ID NO:11, and/or SEQ ID NO:14, or to complements thereof (i.e., SEQ ID NO:16, SEQ ID NO:13, and/or SEQ ID NO:15, respectively).

Oligonucleotides of the present invention can be RNA,

DNA, or derivatives of either. The minimal size of such oligonucleotides is the size required to form a stable hybrid between a given oligonucleotide and the complementary sequence on another nucleic acid molecule of

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Minimal size characteristics are the present invention. disclosed herein. The size of the oligonucleotide must also be sufficient for the use of the oligonucleotide in accordance with the present invention. Oligonucleotides of the present invention can be used in a variety of applications including, but not limited to, as probes to identify additional nucleic acid molecules, as primers to amplify or extend nucleic acid molecules or in therapeutic applications to inhibit, for example, expression of calreticulin proteins by insects. Such therapeutic applications include the use of such oligonucleotides in, for example, antisense-, triplex formation-, ribozymeand/or RNA drug-based technologies. The present invention, therefore, includes such oligonucleotides and methods to interfere with the production of hematophagous insect calreticulin proteins by use of one or more of such technologies.

The present invention also includes a recombinant vector, which includes a hematophagous insect calreticulin nucleic acid molecule of the present invention inserted into any vector capable of delivering the nucleic acid molecule into a host cell. Such a vector contains heterologous nucleic acid sequences, that is nucleic acid sequences that are not naturally found adjacent to a hematophagous insect calreticulin nucleic acid molecule of the present invention. The vector can be either RNA or DNA, either prokaryotic or eukaryotic, and typically is a virus or a plasmid. Recombinant vectors can be used in the

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cloning, sequencing, and/or otherwise manipulating of a hematophagous insect calreticulin nucleic acid molecule of the present invention. One type of recombinant vector, herein referred to as a recombinant molecule and described in more detail below, can be used in the expression of nucleic acid molecules of the present invention. Preferred recombinant vectors are capable of replicating in the transformed cell. Preferred nucleic acid molecules to include in recombinant vectors of the present invention are disclosed herein.

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As heretofore disclosed, one embodiment of the present invention is a method to produce a hematophagous insect calreticulin protein of the present invention by culturing a cell capable of expressing the protein under conditions effective to produce the protein, and recovering the protein. A preferred cell to culture is a recombinant cell that is capable of expressing the hematophagous insect calreticulin protein, the recombinant cell being produced by transforming a host cell with one or more nucleic acid molecules of the present invention. Transformation of a nucleic acid molecule into a cell can be accomplished by any method by which a nucleic acid molecule can be inserted into the cell. Transformation techniques include, but are not limited transfection. electroporation, microinjection, lipofection, adsorption, and protoplast fusion. A recombinant cell may remain unicellular or may grow into a tissue, organ or a multicellular organism. Transformed nucleic acid molecules of the present invention

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can remain extrachromosomal or can integrate into one or more sites within a chromosome of the transformed (i.e., recombinant) cell in such a manner that their ability to be expressed is retained. Preferred nucleic acid molecules with which to transform a host cell are disclosed herein.

Suitable host cells to transform include any cell that can be transformed and that can express the introduced hematophagous insect calreticulin protein. Such cells are, therefore, capable of producing a hematophagous insect calreticulin protein of the present invention after being transformed with at least one nucleic acid molecule of the present invention. Host cells can be either untransformed cells or cells that are already transformed with at least Suitable host cells of the one nucleic acid molecule. present invention can include bacterial, fungal (including. yeast), insect, animal and plant cells. Preferred host cells include bacterial, yeast, insect and mammalian cells, with bacterial (e.g., E. coli) and insect Spodoptera) cells being particularly preferred.

A recombinant cell is preferably produced by transforming a host cell with one or more recombinant molecules, each comprising one or more nucleic acid molecules of the present invention operatively linked to an expression vector containing one or more transcription control sequences. The phrase operatively linked refers to insertion of a nucleic acid molecule into an expression vector in a manner such that the molecule is able to be expressed when transformed into a host cell. As used

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herein, an expression vector is a DNA or RNA vector that is capable of transforming a host cell and of effecting expression of a specified nucleic acid Preferably, the expression vector is also capable of replicating within the host cell. Expression vectors can be either prokaryotic or eukaryotic, and are typically viruses or plasmids. Expression vectors of the present invention include any vectors that function (i.e., direct gene expression) in recombinant cells of the present invention, including in bacterial, fungal, insect, animal, and/or plant cells. As such, nucleic acid molecules of the present invention can be operatively linked to expression vectors containing regulatory sequences such as promoters, operators, repressors, enhancers, termination sequences, origins of replication, and other regulatory sequences that are compatible with the recombinant cell and that control the expression of nucleic acid molecules of the present invention. As used herein, a transcription control sequence includes a sequence which is capable controlling the initiation, elongation, and termination of transcription. Particularly important transcription control sequences are those which control transcription initiation, such as promoter, enhancer, operator and repressor sequences. Suitable transcription control sequences include any transcription control sequence that can function in at least one of the recombinant cells of the present invention. A variety of such transcription control sequences are known to those skilled in the art.

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Preferred transcription control sequences include those which function in bacterial, yeast, helminth, insect and mammalian cells, such as, but not limited to, tac, lac, trp, trc, oxy-pro, omp/lpp, rrnB, bacteriophage lambda (such as lambda p_{μ} and lambda p_{μ} and fusions that include such promoters), bacteriophage T7, T7lac, bacteriophage T3, bacteriophage SP6, bacteriophage SP01, metallothionein, alpha mating factor, Pichia alcohol oxidase, alphavirus subgenomic promoters (such as Sindbis virus subgenomic promoters), baculovirus, Heliothis zea insect virus, vaccinia virus, herpesvirus, poxvirus, adenovirus, simian virus 40, retrovirus actin, retroviral long terminal repeat, Rous sarcoma virus, heat shock, phosphate and nitrate transcription control sequences as well as other sequences capable of controlling gene expression in Additional suitable prokaryotic or eukaryotic cells. transcription control sequences include tissue-specific promoters and enhancers as well as lymphokine-inducible promoters (e.g., promoters inducible by interferons or Transcription control sequences of the interleukins). present invention can also include naturally occurring transcription control sequences naturally associated with a DNA sequence encoding a hematophagous insect calreticulin protein.

Expression vectors of the present invention may also contain secretory signals (i.e., signal segment nucleic acid sequences) to enable an expressed hematophagous insect calreticulin protein to be secreted from the cell that

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produces the protein. Suitable signal segments include a hematophagous insect calreticulin protein signal segment or any heterologous signal segment capable of directing the secretion of a hematophagous insect calreticulin protein, including fusion proteins, of the present invention. Preferred signal segments include, but are not limited to, flea calreticulin, tissue plasminogen activator (t-PA), interferon, interleukin, growth hormone, histocompatibility and viral envelope glycoprotein signal segments.

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Expression vectors of the present invention may also contain fusion sequences which lead to the expression of inserted nucleic acid molecules of the present invention as fusion proteins. Inclusion of a fusion sequence as part of a hematophagous insect calreticulin nucleic acid molecule of the present invention can enhance the stability during . production, storage and/or use of the protein encoded by the nucleic acid molecule. Furthermore, a fusion segment can function as a tool to simplify purification of a hematophagous insect calreticulin protein, such as to enable purification of the resultant fusion protein using affinity chromatography. A suitable fusion segment can be a domain of any size that has the desired function (e.g., increased stability and/or purification tool). within the scope of the present invention to use one or more fusion segments. Fusion segments can be joined to amino and/or carboxyl termini of an insect calreticulin protein. Linkages between fusion segments and hematophagous insect calreticulin proteins be

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constructed to be susceptible to cleavage to enable straight-forward recovery of the hematophagous insect calreticulin proteins. Fusion proteins are preferably produced by culturing a recombinant cell transformed with a fusion nucleic acid sequence that encodes a protein including the fusion segment attached to either the carboxyl and/or amino terminal end of a hematophagous insect calreticulin protein.

A recombinant molecule of the present invention is a molecule that can include at least one of any nucleic acid molecule heretofore described operatively linked to at least one of any transcription control sequence capable of effectively regulating expression of the nucleic acid molecule(s) in the cell to be transformed. A preferred recombinant molecule includes one or more nucleic acid molecules that encode one or more hematophagous insect calreticulin proteins, such as those disclosed herein.

It may be appreciated by one skilled in the art that use of recombinant DNA technologies can improve expression of transformed nucleic acid molecules by manipulating, for example, the number of copies of the nucleic acid molecules within a host cell, the efficiency with which those nucleic acid molecules are transcribed, the efficiency with which the resultant transcripts are translated, and the efficiency of post-translational modifications. Recombinant techniques useful for increasing the expression of nucleic acid molecules of the present invention include, but are not limited to, operatively linking nucleic acid

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molecules to high-copy number plasmids, integration of the nucleic acid molecules into one or more host cell chromosomes, addition of vector stability sequences to plasmids, substitutions or modifications of transcription control signals (e.g., promoters, operators, enhancers), substitutions or modifications of translational control signals (e.g., ribosome binding sites, Shine-Dalgarno sequences), modification of nucleic acid molecules of the present invention to correspond to the codon usage of the host cell, deletion of sequences that destabilize transcripts, and use of control signals that temporally separate recombinant cell growth from recombinant protein production during fermentation. The activity of an expressed recombinant protein of the present invention may be improved by fragmenting, modifying, or derivatizing the resultant protein.

In accordance with the present invention, a recombinant cell can be used to produce a hematophagous insect calreticulin protein of the present invention by culturing such cell under conditions effective to produce such a protein, and recovering the protein. Effective conditions to produce a protein include, but are not limited to, appropriate media, bioreactor, temperature, pH and oxygen conditions that permit protein production. An appropriate, or effective, medium refers to any medium in which a recombinant cell of the present invention, when cultured, is capable of producing a hematophagous insect calreticulin protein. Such a medium is typically an

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aqueous medium comprising assimilable carbohydrate, nitrogen and phosphate sources, as well as appropriate salts, minerals, metals and other nutrients, such as vitamins. The medium may comprise complex nutrients or may be a defined minimal medium.

Cells of the present invention can be cultured in conventional fermentation bioreactors, which include, but are not limited to, batch, fed-batch, cell recycle, and continuous fermentors. Culturing can also be conducted in shake flasks, test tubes, microtiter dishes, and petri plates. Culturing is carried out at a temperature, pH and oxygen content appropriate for the recombinant cell. Such culturing conditions are well within the expertise of one of ordinary skill in the art.

Depending on the vector and host system used for production, resultant hematophagous insect calreticulin proteins may either remain within the recombinant cell; be secreted into the fermentation medium; be secreted into a space between two cellular membranes, such as the periplasmic space in E. coli; or be retained on the outer surface of a cell or viral membrane.

The present invention also includes isolated antibodies against hematophagous insect calreticulin proteins of the present invention (i.e., anti-hematophagous insect calreticulin protein antibodies) and their use to reduce insect infestation on a host animal as well as in the environment of the animal. An anti-hematophagous insect calreticulin protein antibody is an antibody capable

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of selectively binding to a hematophagous insect calreticulin protein. An anti-hematophagous insect calreticulin protein antibody preferably binds to the hematophagous insect calreticulin protein in such a way as to reduce the activity of that protein.

Isolated antibodies are antibodies that have been removed from their natural milieu. The term "isolated" does not refer to the state of purity of such antibodies. such, isolated antibodies can include anti-sera containing such antibodies, or antibodies that have been purified to varying degrees. As used herein, the term "selectively binds to" refers to the ability of such antibodies to preferentially bind to the hematophagous insect calreticulin protein against which the antibody was raised (i.e., to be able to distinguish that protein from unrelated components in a mixture.). Binding affinities typically range from about 103 M.1 to about 1012 M.1. Binding can be measured using a variety of methods known to those skilled in the art including immunoblot assays, immunoprecipitation assays, radioimmunoassays, enzyme immunoassays (e.g., ELISA), immunofluorescent antibody assays and immunoelectron microscopy; see, for example, Sambrook et al., ibid.

Antibodies of the present invention can be either polyclonal or monoclonal antibodies. Antibodies of the present invention include functional equivalents such as antibody fragments and genetically-engineered antibodies, including single chain antibodies, that are capable of

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selectively binding to at least one of the epitopes of the protein used to obtain the antibodies. Antibodies of the present invention also include chimeric antibodies that can bind to more than one epitope. Preferred antibodies are raised in response to proteins that are encoded, at least in part, by a hematophagous insect calreticulin nucleic acid molecule of the present invention.

Anti-hematophagous calreticulin insect antibodies of the present invention include antibodies raised in an animal administered a hematophagous insect calreticulin protein of the present invention that exert their effect when insects feed from the treated animal's blood containing such antibodies. Anti-hematophagous insect calreticulin protein antibodies of the present invention also include antibodies raised in an animal against one or more hematophagous insect calreticulin proteins of the present invention that are then recovered from the animal using techniques known to those skilled in Yet additional antibodies of the present the art. invention are produced recombinantly using techniques as heretofore disclosed for hematophagous insect calreticulin protein of the present invention. Antibodies produced against defined proteins can be advantageous because such antibodies are not substantially contaminated with antibodies against other substances that might otherwise cause interference in a diagnostic assay or side effects if used in a therapeutic composition.

Anti-hematophagous insect calreticulin protein

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antibodies of the present invention have a variety of uses that are within the scope of the present invention. example, such antibodies can be used in a therapeutic composition of the present invention to passively immunize an animal in order to protect the animal from insect Anti-hematophagous insect calreticulin protein antibodies can also be used as tools to screen expression libraries and/or to recover desired proteins of the present invention from a mixture of proteins and other contaminants. Furthermore, antibodies of the present invention can be used to target cytotoxic agents to insects in order to kill insects. Targeting can be accomplished by conjugating (i.e., stably joining) such antibodies to cytotoxic agents using techniques known to those skilled in the art.

A preferred anti-hematophagous insect calreticulin protein antibody of the present invention can selectively bind to, and preferentially reduce calreticulin activity of, a flea calreticulin. A preferred anti-flea calreticulin antibody can selectively bind to a flea calreticulin comprising SEQ ID NO:2 and/or SEQ ID NO:12. A more preferred anti-flea calreticulin antibodies can selectively bind to a flea calreticulin encoded by a nucleic acid molecule comprising SEQ ID NO:1, SEQ ID NO:11 and/or SEQ ID NO:14.

The present invention also includes the use of active calreticulin proteins of the present invention to identify compounds that inhibit calreticulin activity (i.e.,

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calreticulin inhibitory compounds), and preferably a calreticulin inhibitory compound that can be included in a therapeutic composition of the present invention to be administered to animals. A method to identify a compound capable of inhibiting hematophagous insect calreticulin (1) contacting (e.g., combining, activity, comprises: mixing) an isolated hematophagous insect calreticulin protein with a putative (i.e., candidate) inhibitory compound under conditions in which, in the absence of the compound, the protein has calreticulin activity; and (2) determining if the putative inhibitory compound inhibits the calreticulin activity. Putative inhibitory compounds include natural or synthetic molecules, to screen antibodies (including functional equivalents thereof) and Methods to determine calreticulin substrate analogs. activity are known to those skilled in the art. Examples of such methods include measuring the ability of the protein (a) to bind calcium, magnesium, cellular integrins, extracellular matrix proteins, DNA domains of a nuclear hormone receptor, and/or resident endoplasmic reticulum proteins; (b) to affect protein translocation from an endoplasmic reticulum; and/or (c) to act as an agonist or an antagonist of other calcium binding proteins (e.g., calmodulin, calnexin).

The present invention also includes a test kit to identify a compound capable of inhibiting hematophagous insect calreticulin activity. Such a test kit includes an isolated hematophagous insect calreticulin protein having

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calreticulin activity and a means for determining the extent of inhibition of calreticulin activity in the presence of (i.e., effected by) a putative inhibitory compound. Examples of such means are well known to those skilled in the art, some of which are disclosed herein.

The present invention also includes hematophagous insect calreticulin inhibitory compounds isolated by such a method, and/or test kit, and their use to inhibit any hematophagous insect calreticulin that is susceptible to such an inhibitory compound.

It is to be appreciated that the present invention also includes mimetopes of compounds of the present invention that can be used in accordance with methods as disclosed for compounds of the present invention. As used herein, a mimetope of a proteinaceous compound of the present invention (e.g., a hematophagous insect calreticulin protein, an anti-calreticulin antibody, a proteinaceous inhibitor of hematophagous calreticulin activity) refers to any compound that is able to mimic the activity of that proteinaceous compound, often because the mimetope has a structure that mimics that of the proteinaceous compound. For example, a mimetope of a hematophagous insect calreticulin protein is a compound that has an activity similar to that of an isolated hematophagous insect calreticulin protein of the present Mimetopes can be, but are not limited to: peptides that have been modified to decrease their susceptibility to degradation; anti-idiotypic

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catalytic antibodies, or fragments thereof; proteinaceous immunogenic portions of an isolated protein (e.g., carbohydrate structures); and synthetic or natural nucleic organic molecules, including acids Such mimetopes can be designed using carbohydrates. computer-generated structures of proteins of the present invention. Mimetopes can also be obtained by generating random samples of molecules, such as oligonucleotides, peptides or other organic molecules, and screening such samples by affinity chromatography techniques using the corresponding binding partner.

includes therapeutic The present invention compositions, also referred to herein as compositions, that include a (i.e., at least one) compound of the present invention. Preferred compounds to include in a composition of the present invention include an isolated hematophagous insect calreticulin protein or a mimetope thereof, an isolated nucleic acid molecule capable of hybridizing under stringent hybridization conditions with a flea calreticulin gene, an isolated anti-hematophagous insect calreticulin antibody or a mimetope thereof, and/or a calreticulin inhibitory compound, in which the composition, administered to an animal, is able to reduce hematophagous insect burden on the treated animal and/or in the environment of the animal. Such a therapeutic composition can protect an animal from hematophagous insect infestation by reducing calreticulin activity in hematophagous insects animal, thereby reducing feeding from the treated

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hematophagous insect burden on the animal and in the environment of the animal. Suitable and preferred hematophagous insects to target are disclosed herein. Particularly preferred hematophagous insects to target are fleas.

Compositions of the present invention can also include other components such as a pharmaceutically acceptable excipient, an adjuvant, and/or a carrier. For example, compositions of the present invention can be formulated in an excipient that the animal to be treated can tolerate. Examples of such excipients include water, saline, Ringer's solution, dextrose solution, Hank's solution, and other aqueous physiologically balanced salt solutions. Nonaqueous vehicles, such as fixed oils, sesame oil, ethyl oleate, or triglycerides may also be used. Other useful formulations include suspensions containing viscosity enhancing agents, such as sodium carboxymethylcellulose, sorbitol, or dextran. Excipients can also contain minor amounts of additives, such as substances that enhance isotonicity and chemical stability. Examples of buffers include phosphate buffer, bicarbonate buffer and Tris buffer, while examples of preservatives include thimerosal, m- or o-cresol, formalin and benzyl alcohol. formulations can either be liquid injectables or solids which can be taken up in a suitable liquid as a suspension or solution for injection. Thus, in a non-liquid formulation, the excipient can comprise dextrose, human serum albumin, preservatives, etc., to which sterile water

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or saline can be added prior to administration.

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In one embodiment of the present invention, the composition can also include an immunopotentiator, such as an adjuvant or a carrier. Adjuvants are typically substances that generally enhance the immune response of an animal to a specific antigen. Suitable adjuvants include, but are not limited to, Freund's adjuvant; other bacterial cell wall components; aluminum-based salts; calcium-based salts; silica; polynucleotides; toxoids; serum proteins; viral coat proteins; other bacterial-derived preparations; gamma interferon; block copolymer adjuvants, such as Hunter's Titermax™ adjuvant (Vaxcel™, Inc. Norcross, GA); Ribi adjuvants (available from Ribi ImmunoChem Research, Inc., Hamilton, MT); and saponins and their derivatives, such as Quil A (available from Superfos Biosector A/S, Denmark). Carriers are typically compounds that increase the half-life of a therapeutic composition in the treated animal. Suitable carriers include, but are not limited to, polymeric controlled release formulations, biodegradable implants, liposomes, bacteria, viruses, oils, esters, and glycols. One embodiment is an encapsulated therapeutic composition that can be delivered orally.

One embodiment of the present invention is a controlled release formulation that is capable of slowly releasing a composition of the present invention into an animal. As used herein, a controlled release formulation comprises a composition of the present invention in a controlled release vehicle. Suitable controlled release

vehicles include, but are not limited to, biocompatible polymers, other polymeric matrices, capsules, microcapsules, microparticles, bolus preparations, osmotic pumps, diffusion devices, liposomes, lipospheres, and transdermal delivery systems. Other controlled release formulations of the present invention include liquids that, upon administration to an animal, form a solid or a gel in situ. Preferred controlled release formulations are biodegradable (i.e., bioerodible).

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10 A preferred controlled release formulation of the present invention is capable of releasing a composition of the present invention into the blood of the treated animal at a constant rate sufficient to attain therapeutic dose levels of the composition to reduce calreticulin activity 15 in insects feeding from the animal over a period of time ranging from about 1 to about 12 months. A controlled release formulation of the present invention is capable of effecting a treatment for preferably at least about 1 month, more preferably at least about 3 months and even 20 more preferably for at least about 6 months, even more preferably for at least about 9 months, and even more preferably for at least about 12 months.

In order to protect an animal from hematophagous insect infestation, a therapeutic composition of the present invention is administered to the animal in an effective manner such that the calreticulin activity of hematophagous insects feeding from the blood stream of animals treated with the composition is reduced. As such,

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a treated animal is an animal that is competent to reduce hematophagous insect burden by reducing calreticulin activity in insects feeding on that animal. Preferably, the calreticulin activity in the feeding insect is reduced by at least about 50 percent, more preferably by at least about 75 percent, and even more preferably by at least about 100 percent. Methods to administer compositions to the animal in order to render the animal competent depend on the nature of the composition and administration regime.

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Animals administered a composition usually become competent, for example, about 3 to 6 weeks after a primary dose and within another about 2 to 4 weeks with at least one booster shot. Animals administered a composition including anti-hematophagous an insect antibody or calreticulin inhibitory compound become competent as soon as appropriate serum levels of the antibody or inhibitory compound are achieved, usually within about one to about seven days. Methods to measure animal competency (i.e., ability of blood taken up by an insect feeding from the treated animal to reduce calreticulin activity in the insect) are known to those skilled in the art.

In accordance with the present invention, compositions are administered to an animal in a manner such that the animal becomes competent to reduce hematophagous insect calreticulin activity in a hematophagous insect that feeds from the competent animal. For example, a hematophagous insect calreticulin protein of the present invention, when

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administered to an animal in an effective manner (i.e., using an appropriate protocol), is able to elicit (i.e., stimulate) an immune response that produces an antibody titer in the blood stream of the animal sufficient to reduce hematophagous insect calreticulin Similarly, an anti-hematophagous insect calreticulin antibody of the present invention, when administered to an animal in an effective manner, is administered according to a protocol so as to be present in the animal's blood stream at a titer that is sufficient to reduce hematophagous insect calreticulin activity. A calreticulin inhibitory compound of the present invention, when administered to an animal in an effective manner, is administered according to a protocol so as to be present in the animal's blood stream at a is concentration that sufficient to hematophagous insect calreticulin activity. A nucleic acid molecule of the present invention, such as an oligonucleotide or a nucleic acid molecule that encodes a calreticulin protein of the present invention, can also be administered in an effective manner, thereby reducing hematophagous insect calreticulin activity.

Compositions of the present invention can be administered to animals prior to or during hematophagous insect infestation. As noted above, when a hematophagous insect calreticulin protein of the present invention is administered to an animal, a time period is required for the animal to elicit an immune response before the animal is competent to inhibit calreticulin activity of

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hematophagous insects feeding from that animal. Methods to obtain an immune response in an animal are known to those skilled in the art.

Acceptable protocols to administer compositions in an effective manner include individual dose size, number of doses, frequency of dose administration, and mode of Determination of such protocols can be administration. accomplished by those skilled in the art. A suitable single dose is a dose that is capable of protecting an infestation hematophagous insect animal from administered one or more times over a suitable time period. For example, a preferred single dose of calreticulin protein or a mimetope thereof ranges from about 1 microgram (µq) to about 10 milligrams (mg) per kilogram (kg) body Booster vaccinations can be weight of the animal. administered from about 2 weeks to several years after the original administration. Booster immunizations preferably are administered when the immune response of the animal animal from becomes insufficient to protect the Α preferred hematophagous insect infestation. administration schedule is one in which from about 1 μg to about 10 mg of hematophagous insect calreticulin protein per kg body weight of the animal is administered from about one to about two times over a time period of from about 2 weeks to about 12 months. Modes of administration can include, but are not limited to, oral, nasal, topical, transdermal, rectal, and parenteral routes. routes can include, but are not limited to subcutaneous,

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intradermal, intravenous, and intramuscular routes.

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In another embodiment, an anti-hematophagous insect calreticulin antibody composition or a mimetope thereof can be administered in an amount effective to induce a therapeutic effect such as can be determined by one skilled in the art. Anti-hematophagous insect calreticulin antibodies can be re-administered from about 1 hour to about biweekly for several weeks following the original administration. Booster treatments preferably 10 administered when the titer of antibodies of the animal becomes insufficient to protect the animal from hematophagous insect infestation. Suitable modes of administration are as disclosed herein and are known to those skilled in the art.

15 According to one embodiment, a nucleic acid molecule of the present invention can be administered to an animal in a fashion to enable expression of that nucleic acid molecule into a protective protein (e.g., hematophagous insect calreticulin protein, anti-hematophagous insect calreticulin antibody, or proteinaceous calreticulin 20 inhibitory compound) or protective RNA (e.g., antisense RNA, ribozyme or RNA drug) in the animal to be protected from disease. Nucleic acid molecules can be delivered to an animal in a variety of methods including, but not 25 limited to, (a) direct injection (e.g., as "naked" DNA or RNA molecules, such as is taught, for example in Wolff et al., 1990, Science 247, 1465-1468) or (b) packaged as a recombinant virus particle vaccine or as a recombinant cell

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vaccine (i.e., delivered to a cell by a vehicle selected from the group consisting of a recombinant virus particle vaccine and a recombinant cell vaccine).

A recombinant virus particle vaccine of the present invention includes a recombinant molecule of the present invention that is packaged in a viral coat and that can be expressed in an animal after administration. Preferably, the recombinant molecule is packaging-deficient. A number of recombinant virus particles can be used, including, but not limited to, those based on alphaviruses, poxviruses, adenoviruses, herpesviruses, and retroviruses.

When administered to an animal, a recombinant virus particle vaccine of the present invention infects cells within the immunized animal and directs the production of a protective protein or RNA nucleic acid molecule that is capable of protecting the animal from disease caused by the bites of insects of the present invention. A preferred single dose of a recombinant virus particle vaccine of the present invention is from about 1 x 10⁴ to about 1 x 10⁷ virus plaque forming units (pfu) per kilogram body weight of the animal. Administration protocols are similar to those described herein for protein-based vaccines.

A recombinant cell vaccine of the present invention includes recombinant cells of the present invention that express at least one protein of the present invention. Preferred recombinant cells for use in this embodiment include Salmonella, E. coli, Mycobacterium, S. frugiperda, baby hamster kidney, myoblast G8, COS, MDCK and CRFK

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recombinant cells, with Salmonella recombinant cells being more preferred. Such recombinant cells can be administered in a variety of ways but have the advantage that they can be administered orally, preferably at doses ranging from about 10⁸ to about 10¹² bacteria per kilogram body weight. Administration protocols are similar to those described herein for proteinaceous compositions of present invention. Recombinant cell vaccines can comprise whole cells or cell lysates.

Compositions of the present invention can be administered to any animal susceptible to hematophagous insect infestation, including warm-blooded animals. Preferred animals to treat include mammals and birds, with cats, dogs, humans, cattle, chinchillas, ferrets, goats, mice, minks, rabbits, raccoons, rats, sheep, squirrels, swine, chickens, ostriches, quail and turkeys as well as other furry animals, pets and/or economic food animals, being more preferred. Particularly preferred animals to protect are cats, dogs, sheep, cows, pigs, horses and goats.

One embodiment of the present invention is a method to protect an animal from insect infestation, comprising treating an animal with a composition of the present invention. It is to be noted that the term "a" or "an" entity refers to one or more of that entity; for example, a compound refers to one or more compounds. As such, the terms "a" (or "an"), "one or more" and "at least one" can be used interchangeably herein. Thus, a composition of the

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present invention can include one or more compounds that reduce the activity of hematophagous insect calreticulins. Suitable and preferred therapeutic compositions are disclosed herein.

Another embodiment of the present invention includes a method to desensitize a host animal to allergic dermatitis, comprising administering to the animal a hematophagous insect calreticulin protein of the present invention, or a nucleic acid molecule encoding such a protein. Suitable and preferred proteins and nucleic acid molecules are as disclosed for therapeutic compositions useful in reducing hematophagous insect burden in animals. Methods to administer such compounds are as disclosed in U.S. Patent Application Serial No. 08/319,590 (also referred to herein as S/N 08/318,590), ibid.

A hematophagous insect calreticulin protein of the present invention can be used in conjunction with other compounds capable of modifying an animal's hypersensitivity to insect bites. Examples of suitable hematophagous insect saliva proteins to utilize are disclosed in S/N 08/318,590, For example, in order to treat flea allergic ibid. animal can be administered a dermatitis, an present invention calreticulin protein of the combination with one or more flea saliva proteins disclosed in S/N 08/318,590, ibid., including FS-1 flea saliva extract, FS-2 flea saliva extract, and/or proteins included FS-1 and FS-2 flea saliva extracts in those extracts. include mixtures of flea saliva proteins that are collected

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by the method described in detail in S/N 08/319,590, ibid. Preferred flea saliva proteins for use with a flea insect calreticulin protein of the present invention to treat flea allergic dermatitis include at least a portion of one or more of the following flea saliva proteins fspA, fspB, fspC1, fspC2, fspD1, fspD2, fspE, fspF, fspG, fspH, fspI, fspJ1, fspJ2, fspK, fspL1, fspL2, fspM1, fspM2, fspN1, fspN2 and fspN3, as disclosed in S/N 08/319,590, ibid. As used herein and in S/N 08/319,590, ibid., at least a portion of a saliva protein refers to a portion of a saliva protein encoded by a nucleic acid molecule that is capable of hybridizing, under stringent conditions, with a nucleic acid encoding a full-length saliva protein of the invention disclosed in S/N 08,319,590, ibid. More preferred flea saliva proteins for use with a flea calreticulin protein of the present invention include at least a portion of fspE, fspF, fspG, fspH, fspI, fspJ1, fspJ2, fspK, fspL1, fspL2, fspM1, fspM2, fspN1, fspN2 and/or fspN3, with fspG, fspH, fspM1, fspM2, fspN1, fspN2, fspN3, and/or proteins in FS-2 being even more preferred.

Other useful compounds to administer with a hematophagous insect calreticulin protein of the present invention in order to treat allergic dermatitis are compounds that are capable of modifying the function of a cell involved in a hypersensitive response, compounds that reduce allergic reactions, such as systemic agents or anti-inflammatory agents (e.g., anti-histamines, anti-steroid reagents, anti-inflammatory reagents and reagents that

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drive immunoglobulin heavy chain class switching from IgE Suitable compounds useful for modifying the to IqG). function of a cell involved in a hypersensitive response include, but are not limited to, antihistamines, cromolyn sodium, theophylline, cyclosporin A, adrenalin, cortisone, compounds capable of regulating cellular signal transduction, compounds capable of regulating cyclic adenosine monophosphate (cAMP) activity, and compounds that block IgE activity, such as peptides from IgE or IgE specific Fc receptors, antibodies specific for peptides from IgE or IgE-specific Fc receptors, or antibodies capable of blocking binding of IgE to Fc receptors.

Also included in the present invention is the use of a hematophagous insect calreticulin protein of the present invention to identify animals susceptible to or having allergic dermatitis. As used herein, an animal that is susceptible to allergic dermatitis refers to an animal that genetically pre-disposed to developing allergic dermatitis and/or to an animal that has been primed with an antigen in such a manner that re-exposure to the antigen results in symptoms of allergic dermatitis that can be perceived by, for example, observing the animal or measuring antibody production by the animal to the antigen. As such, animals susceptible to allergic dermatitis can include animals having sub-clinical allergic dermatitis. Sub-clinical allergic dermatitis refers to a condition in which allergy symptoms cannot be detected by simply observing an animal, but can be manifest by the presence of

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anti-hematophagous insect calreticulin protein antibodies within an affected animal. Sub-clinical allergic dermatitis can be detected using in vivo or in vitro assays of the present invention as disclosed in S/N 08/319,590, ibid.). Reference to animals having allergic dermatitis includes animals that do display allergy symptoms that can be detected by simply observing an animal and/or by using in vivo or in vitro assays of the present invention.

The following examples are provided for the purposes

of illustration and are not intended to limit the scope of the present invention.

EXAMPLES

Example 1

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This example describes the cloning and sequencing of

a flea calreticulin nucleic acid molecule of the present
invention as well as the deduced amino acid sequence of a
flea calreticulin protein of the present invention.

A flea calreticulin nucleic acid molecule of about 257 nucleotides, referred to herein as nCtCal₂₅₇, was PCR amplified from a flea (Ctenocephalides felis) salivary gland cDNA library that was prepared from RNA isolated from flea salivary glands using standard protocols as described in Sambrook et al., ibid. Primers used in the PCR amplification were degenerate oligonucleotides designed to correspond to conserved regions of the published sequences of human, mouse, rabbit, Onchocerca volvulus and Schistosoma mansoni calreticulin genes, and were as follows: a degenerate "sense" primer having SEQ ID NO:3,

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namely 5'-AARCCNGARGAYTGGGAYAARCC-3' (R is a mixture of A and G; N is a mixture of G, A, T and C; Y is a mixture of C and T); and a degenerate "antisense" primer having SEQ ID NO:4, namely 5'-GTRTAYTCNGGRTTRTCDATYTC-3' (D is a mixture of G, A, and T).

Resulting PCR products were screened using a ³²P-labelled oligonucleotide probe having SEQ ID NO:5, namely 5'- AAYCCNGARGGNGARTGGAA-3', which corresponds to amino acid residues 251 to 258 of the known human calreticulin protein sequence in a standard Southern blot procedure. PCR products that hybridized to the probe were gel purified, cloned into the TA Vector® System (available from Invitrogen, San Diego, CA), and subjected to standard DNA sequencing techniques. An about 257 nucleotide sequence of nCtCal₂₅₇ was determined and is presented as SEQ ID NO:1 (coding strand) and SEQ ID NO:16 (complementary strand). SEQ ID NO:1 apparently encodes a protein of about 85 amino acids, denoted herein as PCtCal₈₅, the amino acid sequence of which is presented as SEQ ID NO:2.

Flea calreticulin nucleic acid sequence SEQ ID NO:1 and flea calreticulin amino acid sequence SEQ ID NO:2 were compared with calreticulin nucleic acid and amino acid sequences characterized from other organisms. SEQ ID NO:1 was found to be about 69% identical to the nucleic acid sequence of the corresponding region of the gene encoding rabbit calreticulin; about 70% identical to the nucleic acid sequence of the corresponding region of the gene encoding human calreticulin; about 71% identical to the

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nucleic acid sequence of the corresponding region of the gene encoding S. mansoni calreticulin; about 74% identical to the nucleic acid sequences of the corresponding regions of the genes encoding mouse calreticulin and rat calreticulin; and about 75% identical to the nucleic acid sequences of the corresponding regions of the genes encoding bovine calreticulin and O. volvulus calreticulin.

SEQ ID NO:2 was found to be about 64% identical to the corresponding region of S. mansoni calreticulin; about 72% identical to the corresponding region of O. volvulus calreticulin; about 81% identical to the corresponding regions of rabbit calreticulin and human calreticulin; and about 82% identical to the corresponding region of mouse calreticulin.

15 Example 2

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This example describes the cloning of additional flea calreticulin nucleic acid molecules of the present invention.

A flea calreticulin nucleic acid molecule of about 850 nucleotides, referred to herein as nCtCal₈₅₀ and including the 5' domain of the flea calreticulin gene, was PCR amplified from a flea salivary gland cDNA library as described in Example 1. The primers used to amplify nCtCal₈₅₀ were as follows: an M13 reverse vector primer having SEQ ID NO:6, namely 5'-GGAAACAGCTATGACCATG-3'; and an antisense primer having SEQ ID NO:7, namely 5'-TGAACCAGACACCTTTGTAGTCAGG-3', which corresponds to nucleotides from about 205 through about 230 of SEQ ID

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NO:1.

Resulting PCR products were screened using a ³²P-labelled oligonucleotide probe corresponding to internal flea calreticulin nucleic acid sequence in a standard Southern blot procedure. PCR products that hybridized to the probe were gel purified, were cloned into the TA Vector® System (available from Invitrogen, San Diego, CA), and were subjected to standard DNA sequencing techniques.

Flea calreticulin nucleic acid molecules of about 550 and of about 556 nucleotides, referred to herein as nCtCalssn and nCtCal₅₅₆, respectively, (each including the 3' domain of the flea calreticulin gene) were PCR amplified from a flea salivary gland cDNA library as described in Example 1. The primers used to amplify the nucleic acid molecules were as follows: an M13 universal vector primer having SEQ ID NO:8, namely 5'-GTAAAACGACGGCCAGT-3'; and a sense primer having SEQ ID NO:9, namely 5'-GGAAGATTGGGACAAGCCAGAAC-3', which corresponds to nucleotides from about 57 through about 79 of SEQ ID NO:1; and another sense primer having SEQ ID NO:10, namely 5'-TTGGGACAAGCCAGAACACATTCC-3', which corresponds to nucleotides from about 63 through about 86 of SEQ ID NO:1. Resulting PCR products were screened using a 32P-labelled oligonucleotide probe having SEQ ID NO:7 in a standard Southern blot procedure. PCR products that hybridized to the probe were gel purified, cloned into the TA Vector® System (available from Invitrogen, San Diego, CA), and subjected to standard DNA sequencing techniques.

Example 3

This Example discloses the production of a recombinant cell of the present invention.

Recombinant molecule pHis-nCtCal₂₅₇, containing flea calreticulin nucleic acid molecule nCtCal₂₅₇ described in Example 1 operatively linked to trc transcription control sequences and to a fusion sequence encoding a polyhistidine segment comprising 6 histidines followed by the amino terminal segment of the bacteriophage S10 capsid protein gene, was produced in the following manner. Nucleic acid molecule nCtCal₂₅₇ was removed from the recombinant vector described in Example 1 by restriction endonuclease digestion, gel purified and subcloned into expression vector pTrcHisA (available from Invitrogen).

The resulting recombinant molecule, denoted pHis-nCtCal₂₅₇, was transformed into *E. coli* to form recombinant cell *E. coli*:pHis-nCtCal₂₅₇.

Example 4

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This Example discloses the production of a flea calreticulin protein of the present invention by a recombinant cell of the present invention.

Recombinant cell *E. coli:*pHis-nCtCal₂₅₇, produced as described in Example 3, is cultured in shake flasks containing an enriched bacterial growth medium containing about 0.1 mg/ml ampicillin at about 37°C. When the cells reach an OD_{600} of about 0.3, expression of $nCtCal_{257}$ is induced by addition of about 1 mM isopropyl- β -D-thiogalactoside (IPTG), and the cells cultured for about 3

hours at about 37°C. Protein production is monitored by SDS PAGE of recombinant cell lysates, followed by Coomassie blue staining, using standard techniques. Recombinant cell E. coli:pHis-nCtCal₂₅₇ produces a fusion protein, denoted herein as PHIS-PCtCal₂₅₇, that migrates with an apparent molecular weight of about 11.5 kD. Such a protein is not produced by cells transformed with the pTrcHisA plasmid lacking a flea nucleic acid molecule insert.

Immunoblot analysis of recombinant cell *E. coli*:pHisnCtCal₂₅₇ lysates indicates that the about 11.5-kD protein binds to a T7 tag monoclonal antibody (available from Novagen, Inc., Madison, WI) directed against the fusion portion of the recombinant PHIS-PCtCal₂₅₇ fusion protein.

Example 5

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This Example describes the cloning of additional flea calreticulin nucleic acid molecules of the present invention.

Based on nucleic acid sequence analysis of nCtCal₈₅₀ and nCtCal₅₅₆, produced as described in Example 2, two additional flea calreticulin gene-specific primers were designed to PCR amplify two additional flea calreticulin nucleic acid molecules from a flea salivary gland cDNA library. Primers used to amplify a flea calreticulin nucleic acid molecule of about 665 nucleotides, referred to herein as nCtCal₆₆₅ and including the 5' domain of the flea calreticulin gene, included: the M13 reverse vector primer having SEQ ID NO:6; and Cal 4R, a flea calreticulin anti-sense primer having SEQ ID NO:17, namely 5' ATTAGGGTCAGGAATAGTTGCACGCTC 3'.

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Primers used to amplify a flea calreticulin nucleic acid molecule of about 750 nucleotides, referred to herein as nCtCal and including the 3' domain of the flea calreticulin gene, included: the M13 universal vector primer having SEQ ID NO:8; and Cal 3F, a flea calreticulin primer having SEQ ID NO:18, namely CATGTATACACTTTGGTTGTTAAGC 3'. Nucleic acid molecules nCtCal and nCtCal were gel purified, cloned into the TA Vector® System (available from Invitrogen), and subjected to standard DNA sequencing techniques.

Yet another flea calreticulin nucleic acid molecule was produced in the following manner. Flea calreticulin nucleic acid molecule nCtCal₁₂₁₈, which includes the entire putative coding region of the flea calreticulin gene, was PCR amplified from a flea salivary gland cDNA library using the following primers: Cal 5F, a flea calreticulin sense primer having SEQ ID NO:19, namely ATAAATATGAAAGCAATTTTGATAACA 3'; and Cal 3R, flea calreticulin anti-sense primer having SEQ ID NO:20, namely 5' TCACAGTTCATCGTGCTCAGCATCGAGTGT 3'. The amplified PCR product was gel purified, cloned into the TA Vector® System (available from Invitrogen), and subjected to standard DNA sequencing techniques.

Nucleic acid sequence analysis of nucleic acid molecules nCtCal₆₆₅, nCtCal₇₅₀, and nCtCal₁₂₁₈ led to the deduction of nucleic acid sequence SEQ ID NO:11 as well as of the complement of SEQ ID NO:11, namely SEQ ID NO:13. As used herein, nucleic acid molecule nCtCal₁₅₈₉ includes SEQ ID

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NO:11 and SEQ ID NO:13. Translation of SEQ ID NO:11 suggests that nucleic acid molecule nCtCal₁₅₈₉ encodes a full-length flea calreticulin protein of about 403 amino acids, referred to herein as PCtCal403, represented by SEQ ID NO:12, assuming an open reading frame having an initiation codon spanning from about nucleotide 151 through 153 and a termination codon spanning from about nucleotide 1360 through 1362 of SEQ ID NO:11. The coding region encoding PCtCal_{ing}, without the stop codon, is represented by nucleic acid molecule nCtCal₁₂₀₀, having the nucleic acid sequence represented by SEQ ID NO:14 (the coding strand) and SEQ ID NO:15 (the complementary strand). The deduced amino acid sequence of SEQ ID NO:12 suggests a protein having a molecular weight of about 46kD and an estimated pI of about 4.23.

Flea calreticulin nucleic acid sequence SEQ ID NO:11 and flea calreticulin amino acid sequence SEQ ID NO:12 were compared with calreticulin nucleic acid and amino acid sequences characterized from other organisms. Since there was very little nucleotide homology in either the 5' or 3' domains between genes of various species, those regions were not compared. SEQ ID NO:11, between about nucleotides 360 and 860, was found to be about 66% identical to the nucleic acid sequence of the corresponding region of the gene encoding bovine calreticulin; about 68% identical to the nucleic acid sequence of the corresponding region of the gene encoding Drosophila calreticulin; about 70% identical the to nucleic acid sequences the

corresponding regions of the genes encoding mouse and rat calreticulins; and about 71% identical to the nucleic acid sequence of the corresponding region of the gene encoding human calreticulin. SEQ ID NO:12, between about amino acids 10 and 360 (a core region least influencedby length or 5' or 3' heterogeneity), was found to be about 68% identical the corresponding region of calreticulin; about 69% identical to the corresponding region of rat calreticulin; about 70% identical to the corresponding regions of mouse calreticulin and human calreticulin; and about 76% identical to the corresponding region of Drosophila calreticulin.

Example 6

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This Example discloses the production of another recombinant cell of the present invention.

Recombinant molecule pTrpHis-nCtCal₁₂₁₂, containing a flea calreticulin nucleic acid molecule comprising the entire putative coding region operatively linked to trp transcription control sequences and to a fusion sequence encoding a poly-histidine segment comprising 6 histidines is produced in the following manner. Nucleic acid molecule nCtCal₁₂₁₂ spanning the flea calreticulin gene from the putative start through stop codons is produced by PCR amplification of the TA Vector® clone containing nCtCal₁₂₁₈ (produced as described in Example 5) using the following primers: Cal-S, a flea calreticulin sense primer having SEQ ID NO:21, namely 5' GAGCTCTCGAGAATAAATATGAAAGCAATTTTG 3'; and Cal-A, a flea calreticulin anti-sense primer having

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SEQ ID NO:22, namely 5' GGACCTCGAGAATCACAGTTCATCGTGCTCAGC 3'. The PCR product is digested with Xho1 restriction endonuclease, gel purified and subcloned into expression vector TrpT2ori/T7-RSET-B (produced as described in PCT Publication No. WO 95/24198, published September 14, 1995, by Tripp et al.), that has been cleaved with Xho1. The resulting recombinant molecule, denoted pTrpHis-nCtCal₁₂₁₂, is transformed into E. coli to form recombinant cell E. coli:pTrpHis-nCtCal₁₂₁₂. Such a recombinant cell, when cultured under effective conditions such as those described in Example 4, leads to the production of a fusion protein denoted herein as PHIS-PCtCal₄₀₃.

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SEQUENCE LISTING

The following Sequence Listing is submitted pursuant to 37 CFR §1.821. A copy in computer readable form is also submitted herewith.

5 Applicants assert pursuant to 37 CFR §1.821(f) that the content of the paper and computer readable copies of SEQ ID NO:1 through SEQ ID NO:22 submitted herewith are the same.

(1) GENERAL INFORMATION:

- 10 (i) APPLICANT: Stiegler, Gary L. Rushlow, Keith E.
 - (ii) TITLE OF INVENTION: HEMATOPHAGOUS INSECT CALRETICULIN NUCLEIC ACID MOLECULES, PROTEINS AND USES THEREOF
- (iii) NUMBER OF SEQUENCES: 22 15
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Sheridan Ross & McIntosh
 - (B) STREET: 1700 Lincoln Street, Suite 3500

 - (C) CITY: Denver (D) STATE: Colorado
 - (E) COUNTRY: U.S.A.
 - (F) ZIP: 80203
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible

 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
- 30 (B) FILING DATE:
 - (C) CLASSIFICATION:
 - (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Connell, Gary J.
 - (B) REGISTRATION NUMBER: 32,020
- 35 (C) REFERENCE/DOCKET NUMBER: 2618-17-C1PCT
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (303) 863-9700
 - (B) TELEFAX: (303) 863-0223

	(2) INFORMATION FOR SEQ ID NO:1:	
5	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 257 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	
10	(ix) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 1255	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
	AAA CCA GAA GAC TGG GAC AAC CGT GCA ACT ATT CCT GAC CCT AAT GAC Lys Pro Glu Asp Trp Asp Asn Arg Ala Thr Ile Pro Asp Pro Asn Asp 1 5 10 15	48
15	ACT AAA CCG GAA GAT TGG GAC AAG CCA GAA CAC ATT CCT GAT CCT GAT Thr Lys Pro Glu Asp Trp Asp Lys Pro Glu His Ile Pro Asp Pro Asp 20 25 30	96
20	GCT ACC AAA CCT GAT GAT TGG GAT GAA GAG ATG GAT GGT GAA TGG GAA Ala Thr Lys Pro Asp Asp Trp Asp Glu Glu Met Asp Gly Glu Trp Glu 35 40 45	144
	CCT GCT ATG ATT GAC AAC CCT GAA TAT AAG GGA GAA TGG GCA CCA AAA Pro Ala Met Ile Asp Asn Pro Glu Tyr Lys Gly Glu Trp Ala Pro Lys 50 55 60	192
25	CAG ATT GAC AAT CCT GAC TAC AAA GGT GTA TGG GTT CAC CCT GAG ATA Gln Ile Asp Asn Pro Asp Tyr Lys Gly Val Trp Val His Pro Glu Ile 65 70 75 80	240
	GAT AAT CCT GAG TAT AC Asp Asn Pro Glu Tyr 85	257

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- (2) INFORMATION FOR SEQ ID NO:2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 85 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Lys Pro Glu Asp Trp Asp Asn Arg Ala Thr Ile Pro Asp Pro Asn Asp 1 5 10 15

Thr Lys Pro Glu Asp Trp Asp Lys Pro Glu His Ile Pro Asp Pro Asp 20 25 30

Ala Thr Lys Pro Asp Asp Trp Asp Glu Glu Met Asp Gly Glu Trp Glu 35 40 45

Pro Ala Met Ile Asp Asn Pro Glu Tyr Lys Gly Glu Trp Ala Pro Lys
50 55 60

Gln Ile Asp Asn Pro Asp Tyr Lys Gly Val Trp Val His Pro Glu Ile 65 70 75 80

Asp Asn Pro Glu Tyr 85

- 20 (2) INFORMATION FOR SEQ ID NO:3:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 23 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 1..23
- 30 (D) OTHER INFORMATION: /label= PRIMER
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

AARCCNGARG AYTGGGAYAA RCC

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	(2) INFOR	RMATION FOR SEQ ID NO:4:	
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	(ii)	MOLECULE TYPE: DNA (genomic)	
10	(ix)	FEATURE: (A) NAME/KEY: misc_feature (B) LOCATION: 123 (D) OTHER INFORMATION: /label= PRIMER	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:4:	
	GTRTAYTC	NG GRTTRTCDAT YTC	2
	(2) INFOR	RMATION FOR SEQ ID NO:5:	
15	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
20	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE: (A) NAME/KEY: misc_feature (B) LOCATION: 120 (D) OTHER INFORMATION: /label= PRIMER	
25	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:5:	
	AAYCCNGAR	RG GNGARTGGAA	20
	(2) INFOR	RMATION FOR SEQ ID NO:6:	
30.	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
35	(ix)	FEATURE: (A) NAME/KEY: misc_feature (B) LOCATION: 119 (D) OTHER INFORMATION: /label= PRIMER	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:6:	
	GGAAACAGC	T ATGACCATG	19

	(2) INFORMATION FOR SEQ ID NO:7:	
5	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
10	<pre>(ix) FEATURE: (A) NAME/KEY: misc feature (B) LOCATION: 125 (D) OTHER INFORMATION: /label= PRIMER</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
	TGAACCAGAC ACCTTTGTAG TCAGG	25
	(2) INFORMATION FOR SEQ ID NO:8:	
15	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
20	(ii) MOLECULE TYPE: DNA (genomic)	
	<pre>(ix) FEATURE: (A) NAME/KEY: misc_feature (B) LOCATION: 117 (D) OTHER INFORMATION: /label= PRIMER</pre>	
25	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
	GTAAAACGAC GGCCAGT	17
	(2) INFORMATION FOR SEQ ID NO:9:	
30	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
35	<pre>(ix) FEATURE: (A) NAME/KEY: misc feature (B) LOCATION: 123 (D) OTHER INFORMATION: /label= PRIMER</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
	GGAAGATTGG GACAAGCCAG AAC	23

	(2) INFORMATION FOR SEQ ID NO:10:	
5	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
10	<pre>(ix) FEATURE: (A) NAME/KEY: misc_feature (B) LOCATION: 124 (D) OTHER INFORMATION: /label= PRIMER</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
	TTGGGACAAG CCAGAACACA TTCC	2
	(2) INFORMATION FOR SEQ ID NO:11:	
15	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 1589 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
20	(ii) MOLECULE TYPE: cDNA	
	(ix) FEATURE: (A) NAME/KEY: CDS (B) LOCATION: 1511360	
) E	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	6
25	GGCACGAGCG ACGACGAATG CATAAGCCGC ATTTGTTATT GAGAAATATA TGCTAAAATA TATTGTCTCA GAATTATTAT AATGTGCTGA GTTTGCATTA ATATTGAAAC GTTTTATTCA	12
	TTCTGAGGCA TCTATATAAA ATAAATAAAT ATG AAA GCA ATT TTG ATA ACA TTG Met Lys Ala Ile Leu Ile Thr Leu 1 5	17
30	ATA GTC GCC GCG GCT GTG TAT TCC GTA AGG CCT GAG GTT TTC CTG GAA Ile Val Ala Ala Ala Val Tyr Ser Val Arg Pro Glu Val Phe Leu Glu 10 15 20	22
35	GAA AAC TTC GTA GAC GAT ACG TGG ACA AAT ACA TGG GTT TAT AGT GAA Glu Asn Phe Val Asp Asp Thr Trp Thr Asn Thr Trp Val Tyr Ser Glu 25 30 35 40	27
٠	CAC CCT GGC AAA GAA TTC GGC AAA TTC GTG CAC ACT GCC GGA AAG TTC His Pro Gly Lys Glu Phe Gly Lys Phe Val His Thr Ala Gly Lys Phe 45 50 55	31
40	TAT AAC GAT GCC GAA GCA GAC AAA GGT TTG CAA ACA GGT CAA GAT GCT Tyr Asn Asp Ala Glu Ala Asp Lys Gly Leu Gln Thr Gly Gln Asp Ala 60 65 70	36
	AGG TTC TAC GCT CTA TCT CAT AAG TTC AAA CCT TTC TCA AAT AAA GAC Arg Phe Tyr Ala Leu Ser His Lys Phe Lys Pro Phe Ser Asn Lys Asp	414

	AA Ly	s Th	A TT r Le O	A GT u Va	T GT	A CAA 1 Gl	n Ph 9	e Se	C GTT er Va	r aas 1 Ly	CAT B Hi	GAA B Gl 10	u Gl	AAC n Asn	ATT GAC Ile As	462 p
5	TG' Cy: 10!	B Gl	A GG y Gl	T GG Y Gl	T TAC y Ty	TTG Lev 110	ı Ly	GG1 B Gl	r TTC y Ph	GAA e Gl	TTC u Ph 11	e Se	GTG r Va	AAT (l Asn	CAA AAG Gln Ly: 120	-
	GA(As)	C ATO	G CA: t Hi	r GG(G GAF Y Gli 12:	u Ser	CCC Pro	TAT O Ty	GAA	ATT u Il 13	e Me	TTT t Pho	GGT e Gly	CCT (SAC ATT Asp Ile 135	558 e
10	TG1 Cys	GAG B Asj	C CCI	A GG/ O G1: 140	y Thi	AAG Lys	AAG Lys	GTT B Va	CAC 1 Hi	в Va	ATC l Ile	TTC e Phe	AGC e Ser	TAC A Tyr 150	AG GGT Lys Gl _y	606
15	AA# Lys	AAA BABI	7 GT1 1 Val 159	l Lei	ATC	AAT Asn	AAG Lys	GAT Asj 160	o Ile	CGC Arg	TGC g Cyt	AAA S Lys	GAT Asp 165	Asp	TC TAT Val Tyr	654
	ACT Thr	CAT His	val	TAC Tyr	ACT Thr	TTG Leu	GTT Val 175	. Val	AAG L Lys	CCC Pro	GAT Asp	AAT Asn 180	Thr	TAT G	AG GTG Glu Val	702
20	TTG Leu 185	Ile	GAT Asp	AAT Asr	GAG	AAG Lys 190	GTT Val	GAA Glu	AGT Ser	GGT Gly	AAC Asn 195	Leu	GAA Glu	GAT G Asp	AC TGG Asp Trp 200	750
	GAA Glu	TTC Phe	CTA Leu	GCC Ala	CCC Pro 205	Lys	AAA Lys	ATC Ile	AAG Lys	GAT Asp 210	Pro	GAA Glu	GCT A	Lys	AA CCA Lys Pro 215	798
25	GCA Ala	GAT Asp	TGG Trp	GAT Asp 220	Glu	CGT Arg	GCA Ala	ACT Thr	ATT Ile 225	Pro	GAC Asp	CCT Pro	AAT (Asp 230	CC AAA Thr Lys	846
30	CCT Pro	GAA Glu	GAT Asp 235	Trp	GAC Asp	AAG Lys	CCA Pro	GAA Glu 240	His	ATT Ile	CCT Pro	GAT (Asp	Pro 245	AT GO	CT ACC Ala Thr	894
	AAA Lys	CCT Pro 250	Asp	GAC Asp	TGG Trp	GAT (GAA Glu 255	GAG Glu	ATG Met	GAT Asp	GGT (GAA 1 Glu 260	ree e Trp	AA CC Glu I	T GCT Pro Ala	942
35	ATG Met 265	ATT Ile	GAC Asp	AAC Asn	CCT Pro	GAA 1 Glu 270	TAT . Tyr	AAG Lys	GGA (GAA ' Glu	TGG (Trp 275	GCA C Ala	CA A Pro	AA CA Lys C	G ATT Sln Ile 280	990
	GAC Asp	AAT Asn	CCT Pro	GAC Asp	TAC Tyr 285	AAA (Lys	GT (GTC ' Val	TGG (Trp	Val 290	CAC (His	CT G Pro	AA A Glu	Ile A	T AAT Asp Asn 95	1038
40	CCA Pro	GAA Glu	TAT Tyr	GTT Val 300	CCT Pro	GAT A	CT (Thr	Gln	CTT 1 Leu 305	TAC A	Lys Lys	GT G Arg	Asp	AG AT Glu I 310	T TGT le Cys	1086
45	GCC Ala	ATT Ile	GGT Gly 315	TTA Leu	GAT '	TTA 1 Leu	Trp	GAA (Gln 320	STA / Val	AG G Lys	CT G Ala	Gly	CA AT Thr 325	TA TTO	C GAC he Asp	1134
	Asn	ATT Ile 330	TTA . Leu	ATC . Ile	ACA (yab y	АТ G Авр 335	TT C Val	ASP	'AT G Tyr	Ala	AG AI Lys 1 340	AA A1 Lys :	TA GC/ Ile A	A GAA la Glu	1182
50	GGT Gly 345	GTT . Val	AAA ' Lys	TCT : Ser	Thr	CAG G Gln (350	AA G Glu (GA G	AA A Glu	Lys	AA A Lys : 355	TG AI Met 1	AA GA Lys <i>I</i>	T GCT	CAA la Gln 360	1230

						Lys		AGG Arg			Glu					1 Asn	1278
5					Asp			TTA Leu		Asp					Pro	GAA O Glu	1326
								CAC His 400	Asp				TTTT	TAAG	}		1370
10	TGC:	TACT	CAC (CATA	AACT:	TT TO	CACAT	TGGC	TTA	ATTI	ATT	TCCG	TTAA	AT C	ATCC	AACAT	1430
	CTATACATTA ATTATTACCT TGTAGAAAAT TGTGTTTGTG AAAAATTTGT CTCCGTTTAC													1490			
	TTGAAACAAT GAAGTGCATG CCAATTGTGT AATAATCGAC TGTGCCCAAA ATAAATTATT													1550			
	TAA!	TTCT'	TGT :	rcaa:	FAA G	AT T	TTGT	TATA	C GT	AAGT'	TTT						1589
	(2)	INF	ORMA'	rion	FOR	SEQ	ID :	NO:1	2:								
15			(i)					ERIS'			a						
				(B	TY!	PE:	amin	o ac line	id	2014							
			i i)	•	•			rote									
20		-	•				_	TION		O ID	NO:	12:					
	Met	•	•	_									Ala	Val	Tvr	Ser	
	1	 ,.			5					10					15		
		-		20					25					30		Trp	
25			35	_				Glu 40					45				
	Phe	Val 50	His	Thr	Ala	Gly	Lys 55	Phe	Tyr	Asn	Asp	Ala 60	Glu	Ala	Asp	Lys	
30	Gly 65	Leu	Gln	Thr	Gly	Gln 70	Asp	Ala	Arg	Phe	Tyr 75	Ala	Leu	Ser	His	Lys 80	
	Phe	Lys	Pro	Phe	Ser 85	Asn	Lys	Asp	Lys	Thr 90	Leu	Val	Val	Gln	Phe 95	Ser	
	Val	Lys	His	Glu 100	Gln	Asn	Ile	yab	Cys 105	Gly	Gly	Gly	Tyr	Leu 110	Lys	Gly	
35	Phe	Glu	Phe 115	Ser	Val	Asn	Gln	Lys 120	Asp	Met	His	Gly	Glu 125	Ser	Pro	Tyr	
		130					135	Ile				140					
40	145					150		Gly			155					160	
	Ile	Arg	Сув	Lys	Asp 165	Asp	Val	Tyr	Thr	His 170	Val	Tyr	Thr	Leu	Val 175	Val	

	Lys	B Pro) Asp) Asn 180	Thr	Туг	Glu	ı Val	Let 185	ı Ile	e Ast) Ası	ı Glı	1 Ly:		l Gl
	Ser	Gly	195	Leu	Glu	Asp) Asp	200	Glu	Phe	Leu	Ala	205		E Ly	s Il
5	Lye	210	Pro	Glu	Ala	Lys	Lys 215	Pro	Ala	Asp	Trp	Asp 220	Glu	Arç	, Ala	a Th
	Ile 225	Pro	Asp	Pro	Asn	Asp 230	Thr	Lys	Pro	Glu	Asp 235	Trp	Asp	Lys	Pro	Gl: 240
10	His	Ile	Pro	Asp	Pro 245	Asp	Ala	Thr	Lys	Pro 250	Asp	Asp	Trp	Asp	Glu 255	
	Met	Asp	Gly	Glu 260	Trp	Glu	Pro	Ala	Met 265	Ile	Asp	Asn	Pro	Glu 270		Lys
	Gly	Glu	Trp 275	Ala	Pro	Lys	Gln	Ile 280	yab	Asn	Pro	Asp	Tyr 285	Lys	Gly	Val
15	Trp	Val 290	His	Pro	Glu	Ile	Asp 295	Asn	Pro	Glu	Tyr	Val 300	Pro	Asp	Thr	Gln
	Leu 305	Tyr	Lys	Arg	Asp	Glu 310	Ile	Сув	Ala	Ile	Gly 315	Leu	Asp	Leu	Trp	Gln 320
20	Val	Lys	Ala	Gly	Thr 325	Ile	Phe	Asp	Asn	Ile 330	Leu	Ile	Thr	Asp	Asp 335	Val
	Asp	Tyr	Ala	Lys 340	Lys	Ile	Ala	Glu	Gly 345	Val	Lys	Ser	Thr	Gln 350	Glu	Gly
	Glu	Lys	Lys 355	Met	Lys	Asp	Ala	Gln 360	Asp	Glu	Glu	Glu	Arg 365	Lys	Ala	Arg
25	Glu	Ala 370	Glu	Thr	Lys	Glu	Glu 375	Asn	Asp	Thr		Ala 380	Asp	Glu	Asp	Leu
	Asp 385	Asp	Asn	Ala	Glu	Thr 390	Pro	Glu	Glu		Thr :	Leu	Asp	Ala	Glu	His 400
	Asp	Glu	Leu													

70

(2) INFORMATION FOR SEQ ID NO:13:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 1589 base pairs
- (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

AAAACTTACG TATAACAAAA TCTTATTGAA CAAGAATTAA ATAATTTATT TTGGGCACAG 60 10 TCGATTATTA CACAATTGGC ATGCACTTCA TTGTTTCAAG TAAACGGAGA CAAATTTTTC 120 ACAAACACAA TTTTCTACAA GGTAATAATT AATGTATAGA TGTTGGATGA TTTAACGGAA 180 ATAAATTAAG CCAATGTGAA AAGTTTATGG TGAGTAGCAC TTAAAAATCA CAGTTCATCG 240 TGCTCAGCAT CGAGTGTGTC TTCTTCTGGT GTTTCGGCAT TATCATCTAA GTCTTCATCA 300 GCATCTGTGT CATTTTCTTC TTTTGTTTCG GCTTCCCTGG CTTTCCTTTC TTCTTCATCT 360 15 TGAGCATCTT TCATTTTCTT TTCTCCTTCC TGGGTAGATT TAACACCTTC TGCTATTTTC 420 TTTGCATAAT CAACATCATC TGTGATTAAA ATATTGTCGA ATATTGTTCC AGCCTTTACT 480 TGCCATAAAT CTAAACCAAT GGCACAAATC TCATCACGTT TGTAAAGTTG AGTATCAGGA 540 ACATATTCTG GATTATCAAT TTCAGGGTGA ACCCAGACAC CTTTGTAGTC AGGATTGTCA 600 ATCTGTTTTG GTGCCCATTC TCCCTTATAT TCAGGGTTGT CAATCATAGC AGGTTCCCAT 660 20 TCACCATCCA TCTCTTCATC CCAGTCATCA GGTTTGGTAG CATCAGGATC AGGAATGTGT 720 780 TCTGGCTTGT CCCAATCTTC AGGTTTGGTG TCATTAGGGT CAGGAATAGT TGCACGCTCA TCCCAATCTG CTGGTTTTTT AGCTTCTGGA TCCTTGATTT TCTTGGGGGC TAGGAATTCC 840 CAGTCATCTT CCAAGTTACC ACTTCAACC TTCTCATTAT CAATCAACAC CTCATAGGTA 900 TTATCGGGCT TAACAACCAA AGTGTATACA TGAGTATAGA CATCATCTTT GCAGCGGATA 960 25 TCCTTATTGA TCAAAACATT TTTACCCTTG TAGCTGAAGA TTACGTGAAC CTTCTTAGTT 1020 CCTGGGTCAC AAATGTCAGG ACCAAACATA ATTTCATAGG GACTTTCCCC ATGCATGTCC 1080 TTTTGATTCA CACTGAATTC GAAACCCTTC AAGTAACCAC CTCCACAGTC AATGTTTTGT 1140 TTATGAGATA GAGCGTAGAA CCTAGCATCT TGACCTGTTT GCAAACCTTT GTCTGCTTCG 1260 30 GCATCGTTAT AGAACTTTCC GGCAGTGTGC ACGAATTTGC CGAATTCTTT GCCAGGGTGT 1320 TCACTATAAA CCCATGTATT TGTCCACGTA TCGTCTACGA AGTTTTCTTC CAGGAAAACC 1380 TCAGGCCTTA CGGAATACAC AGCCGCGGCG ACTATCAATG TTATCAAAAT TGCTTTCATA 1440 TTTATTTATT TTATATAGAT GCCTCAGAAT GAATAAAACG TTTCAATATT AATGCAAACT 1500 CAGCACATTA TAATAATTCT GAGACAATAT ATTTTAGCAT ATATTTCTCA ATAACAAATG 1560 35 1589 CGGCTTATGC ATTCGTCGTC GCTCGTGCC

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(2) INFORMATION FOR SEQ ID NO:14:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 1209 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

ATGAAAGCAA TTTTGATAAC ATTGATAGTC GCCGCGGCTG TGTATTCCGT AAGGCCTGAG 60 10 GTTTTCCTGG AAGAAACTT CGTAGACGAT ACGTGGACAA ATACATGGGT TTATAGTGAA 120 CACCCTGGCA AAGAATTCGG CAAATTCGTG CACACTGCCG GAAAGTTCTA TAACGATGCC 180 GAAGCAGACA AAGGTTTGCA AACAGGTCAA GATGCTAGGT TCTACGCTCT ATCTCATAAG 240 TTCAAACCTT TCTCAAATAA AGACAAGACA TTAGTTGTAC AATTTTCCGT TAAACATGAA 300 CAAAACATTG ACTGTGGAGG TGGTTACTTG AAGGGTTTCG AATTCAGTGT GAATCAAAAG 360 GACATGCATG GGGAAAGTCC CTATGAAATT ATGTTTGGTC CTGACATTTG TGACCCAGGA 420 ACTAAGAAGG TTCACGTAAT CTTCAGCTAC AAGGGTAAAA ATGTTTTGAT CAATAAGGAT 480 ATCCGCTGCA AAGATGATGT CTATACTCAT GTATACACTT TGGTTGTTAA GCCCGATAAT 540 ACCTATGAGG TGTTGATTGA TAATGAGAAG GTTGAAAGTG GTAACTTGGA AGATGACTGG 600 GAATTCCTAG CCCCCAAGAA AATCAAGGAT CCAGAAGCTA AAAAACCAGC AGATTGGGAT 660 GAGCGTGCAA CTATTCCTGA CCCTAATGAC ACCAAACCTG AAGATTGGGA CAAGCCAGAA 720 CACATTCCTG ATCCTGATGC TACCAAACCT GATGACTGGG ATGAAGAGAT GGATGGTGAA 780 TGGGAACCTG CTATGATTGA CAACCCTGAA TATAAGGGAG AATGGGCACC AAAACAGATT 840 GACAATCCTG ACTACAAAGG TGTCTGGGTT CACCCTGAAA TTGATAATCC AGAATATGTT 900 CCTGATACTC AACTTTACAA ACGTGATGAG ATTTGTGCCA TTGGTTTAGA TTTATGGCAA 960 GTAAAGGCTG GAACAATATT CGACAATATT TTAATCACAG ATGATGTTGA TTATGCAAAG 1020 AAAATAGCAG AAGGTGTTAA ATCTACCCAG GAAGGAGAAA AGAAAATGAA AGATGCTCAA 1080 GATGAAGAAG AAAGGAAAGC CAGGGAAGCC GAAACAAAAG AAGAAAATGA CACAGATGCT 1140 GATGAAGACT TAGATGATAA TGCCGAAACA CCAGAAGAAG ACACACTCGA TGCTGAGCAC 1200 GATGAACTG 1209

PCT/US96/03133

72

(2) INFORMATION FOR SEQ ID NO:15:

WO 96/28469

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(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 1209 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

	CAGTTCATCG	TGCTCAGCAT	CGAGTGTGTC	TTCTTCTGGT	GTTTCGGCAT	TATCATCTAA	60
10	GTCTTCATCA	GCATCTGTGT	CATTTTCTTC	TTTTGTTTCG	GCTTCCCTGG	CTTTCCTTTC	120
	TTCTTCATCT	TGAGCATCTT	TCATTTTCTT	TTCTCCTTCC	TGGGTAGATT	TAACACCTTC	180
	TGCTATTTTC	TTTGCATAAT	CAACATCATC	TGTGATTAAA	ATATTGTCGA	ATATTGTTCC	240
	AGCCTTTACT	TGCCATAAAT	CTAAACCAAT	GGCACAAATC	TCATCACGTT	TGTAAAGTTG	300
	AGTATCAGGA	ACATATTCTG	GATTATCAAT	TTCAGGGTGA	ACCCAGACAC	CTTTGTAGTC	360
15	AGGATTGTCA	ATCTGTTTTG	GTGCCCATTC	TCCCTTATAT	TCAGGGTTGT	CAATCATAGC	420
	AGGTTCCCAT	TCACCATCCA	TCTCTTCATC	CCAGTCATCA	GGTTTGGTAG	CATCAGGATC	480
	AGGAATGTGT	TCTGGCTTGT	CCCAATCTTC	AGGTTTGGTG	TCATTAGGGT	CAGGAATAGT	540
	TGCACGCTCA	TCCCAATCTG	CTGGTTTTTT	AGCTTCTGGA	TCCTTGATTT	TCTTGGGGGC	600
	TAGGAATTCC	CAGTCATCTT	CCAAGTTACC	ACTTTCAACC	TTCTCATTAT	CAATCAACAC	660
20	CTCATAGGTA	TTATCGGGCT	TAACAACCAA	AGTGTATACA	TGAGTATAGA	CATCATCTTT	720
	GCAGCGGATA	TCCTTATTGA	TCAAAACATT	TTTACCCTTG	TAGCTGAAGA	TTACGTGAAC	780
	CTTCTTAGTT	CCTGGGTCAC	AAATGTCAGG	ACCAAACATA	ATTTCATAGG	GACTTTCCCC	840
	ATGCATGTCC	TTTTGATTCA	CACTGAATTC	GAAACCCTTC	AAGTAACCAC	CTCCACAGTC	900
	AATGTTTTGT	TCATGTTTAA	CGGAAAATTG	TACAACTAAT	GTCTTGTCTT	TATTTGAGAA	960
25	AGGTTTGAAC	TTATGAGATA	GAGCGTAGAA	CCTAGCATCT	TGACCTGTTT	GCAAACCTTT	1020
	GTCTGCTTCG	GCATCGTTAT	AGAACTTTCC	GGCAGTGTGC	ACGAATTTGC	CGAATTCTTT	1080
	GCCAGGGTGT	TCACTATAAA	CCCATGTATT	TGTCCACGTA	TCGTCTACGA	AGTTTTCTTC	1140
	CAGGAAAACC	TCAGGCCTTA	CGGAATACAC	AGCCGCGGCG	ACTATCAATG	TTATCAAAAT	1200
	TGCTTTCAT						1209

÷	(2) INFORMATION FOR SEQ ID NO:16:	
5	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 257 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:	
	GTATACTCTG GATTATCTAT TTCAGGGTGA ACCCAGATAC CTTTGTAGTC AGGATTGTCA	60
10	ATCTGTTTTG GTGCCCATTC TCCCTTATAT TCAGGGTTGT CAATCATAGC AGGTTCCCAT	120
	TCACCATCCA TCTCTTCATC CCAATCATCA GGTTTGGTAG CATCAGGATC AGGAATGTGT	180
	TCTGGCTTGT CCCAATCTTC CGGTTTAGTG TCATTAGGGT CAGGAATAGT TGCACGGTTG	240
	TCCCAGTCTT CTGGTTT	257
	(2) INFORMATION FOR SEQ ID NO:17:	
15	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
20	(ii) MOLECULE TYPE: DNA (genomic)	
	<pre>(ix) FEATURE: (A) NAME/KEY: misc feature (B) LOCATION: 127 (D) OTHER INFORMATION: /label= primer</pre>	
25	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:	
	ATTAGGGTCA GGAATAGTTG CACGCTC	27
	(2) INFORMATION FOR SEQ ID NO:18:	
3 0	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
35	<pre>(ix) FEATURE: (A) NAME/KEY: misc_feature (B) LOCATION: 125 (D) OTHER INFORMATION: /label= primer</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:	
	CATGTATACA CTTTGGTTGT TAAGC	25

	(2) INFORMATION FOR SEQ ID NO:19:	
5	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
10	<pre>(ix) FEATURE: (A) NAME/KEY: misc_feature (B) LOCATION: 127 (D) OTHER INFORMATION: /label= primer</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:	
	ATAAATATGA AAGCAATTTT GATAACA	27
	(2) INFORMATION FOR SEQ ID NO:20:	
15	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
20	(ii) MOLECULE TYPE: DNA (genomic)	
	<pre>(ix) FEATURE: (A) NAME/KEY: misc_feature (B) LOCATION: 130 (D) OTHER INFORMATION: /label= primer</pre>	
25	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:	
	TCACAGTTCA TCGTGCTCAG CATCGAGTGT	30
	(2) INFORMATION FOR SEQ ID NO:21:	
30	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
35	<pre>(ix) FEATURE: (A) NAME/KEY: misc_feature (B) LOCATION: 133 (D) OTHER INFORMATION: /label= primer</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:	
	GAGCTCTCGA GAATAAATAT GAAAGCAATT TTG	33

75

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- (2) INFORMATION FOR SEQ ID NO:22:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 33 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:

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- (A) NAME/KEY: misc_feature
- 10 (B) LOCATION: 1..33
 - (D) OTHER INFORMATION: /label= primer
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22: GGACCTCGAG AATCACAGTT CATCGTGCTC AGC

While various embodiments of the present invention have been described in detail, it is apparent that 15 modifications and adaptations of those embodiments will occur to those skilled in the art. It is to be expressly understood, however, that such modifications and . adaptations are within the scope of the present invention, 20 as set forth in the following claims.

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What is claimed is:

- 1. An isolated hematophagous insect calreticulin protein encoded by a nucleic acid molecule that hybridizes under stringent hybridization conditions to a flea calreticulin gene.
- 2. The protein of Claim 1, wherein said flea calreticulin gene comprises nucleic acid sequence SEQ ID NO:1.
- 3. The protein of Claim 1, wherein said flea calreticulin gene encodes a protein comprising amino acid sequence SEQ ID NO:2.
 - 4. The protein of Claim 1, wherein said protein can be isolated from insect saliva.
- 5. The protein of Claim 1, wherein said protein comprises an amino acid sequence that is at least about 85% identical to an amino acid sequence represented by an amino acid sequence selected from the group consisting of SEQ ID NO:2 and SEQ ID NO:12.
- 6. The protein of Claim 1, wherein said protein comprises an amino acid sequence selected from the group consisting of SEQ ID NO:2 and SEQ ID NO:12.
 - 7. The protein of Claim 1, wherein said protein is encoded by a nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:11 and SEQ ID NO:14.
 - 8. The protein of Claim 1, wherein said protein, when administered to an animal, is capable of reducing insect infestation by a method comprising altering the

77

blood feeding behavior of insects that feed from said treated animal.

9. The protein of Claim 1, wherein said protein, when administered to an animal, reduces calreticulin activity in insects feeding from said animal, thereby reducing insect burden on said animal and in the environment of said animal.

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- 10. The protein of Claim 1, wherein said protein, when administered to an animal, is capable of eliciting an immune response against an insect calreticulin.
- 11. The protein of Claim 1, wherein said protein, when administered to an animal, is capable of substantially desensitizing said animal to allergic dermatitis.
- 12. The protein of Claim 11, wherein said allergic

 dermatitis is selected from the group consisting of flea

 allergic dermatitis, mosquito allergic dermatitis and

 Culicoides allergic dermatitis.
 - 13. The protein of Claim 1, wherein said protein is used to identify animals susceptible to or having allergic dermatitis.
 - 14. An isolated antibody capable of selectively binding to an isolated protein as set forth in Claim 1.
 - 15. An isolated hematophagous insect calreticulin nucleic acid molecule that hybridizes under stringent hybridization conditions to a flea calreticulin gene.
 - 16. The nucleic acid molecule of Claim 15, wherein said nucleic acid molecule hybridizes under stringent hybridization conditions to a nucleic acid sequence

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so lected from the group consisting of SEQ ID NO:1, SEQ ID NO:11, SEQ ID NO:13, SEQ ID NO:14, SEQ ID NO:15, and SEQ ID NO:16.

- 17. The nucleic acid molecule of Claim 15, wherein said nucleic acid molecule hybridizes under stringent hybridization conditions to a nucleic acid sequence encoding a protein comprising an amino acid sequence selected from the group consisting of SEQ ID NO:2 and SEQ ID NO:12.
- 18. The nucleic acid molecule of Claim 15, wherein said nucleic acid molecule comprises a nucleic acid sequence having at least about 80% identity with nucleic acid sequence SEQ ID NO:1.
 - 19. The nucleic acid molecule of Claim 15, wherein said nucleic acid molecule comprises an oligonucleotide.
 - 20. The nucleic acid molecule of Claim 15, wherein said nucleic acid sequence encodes a hematophagous insect calreticulin protein having calcium binding activity.
- 21. The nucleic acid molecule of Claim 15 or the protein of Claim 1, wherein said insect is selected from the group consisting of fleas, midges, mosquitos, sand flies, black flies, horse flies, horn flies, deer flies, tsetse flies, buffalo flies, blow flies, stable flies, myiasis-causing flies, biting gnats, lice and true bugs.
- 25 22. The nucleic acid molecule of Claim 15 or the protein of Claim 1, wherein said insect is a flea of a genus selected from the group consisting of Ctenocephalides, Cyopsyllus, Diamanus, Echidnophaga,

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Nosopsyllus, Pulex, Tunga, Oropsylla, Orchopeus and Xenopsylla.

- 23. The nucleic acid molecule of Claim 15 or the protein of Claim 1,, wherein said insect is a flea of a species selected from the group consisting of Ctenocephalides felis, Ctenocephalides canis, Pulex irritans, Oropsylla (Thrassis) bacchi, Oropsylla (Diamanus) montana, Orchopeus howardi, Xenopsylla cheopis and Pulex simulans.
- 24. The nucleic acid molecule of Claim 15, wherein said nucleic acid molecule encodes a protein capable of eliciting an immune response against a hematophagous insect calreticulin.
- 25. The nucleic acid molecule of Claim 15, wherein said nucleic acid molecule, when administered to an animal, reduces calreticulin activity in insects feeding from said animal, thereby reducing insect burden on said animal and in the environment of said animal.
- 26. The nucleic acid molecule of Claim 15, wherein said nucleic acid molecule comprises a nucleic acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:11, SEQ ID NO:13, SEQ ID NO:14, SEQ ID NO:15, and SEQ ID NO:16.
- 27. The nucleic acid molecule of Claim 15, wherein said nucleic acid molecule encodes a protein comprising an amino acid sequence selected from the group consisting of SEQ ID NO:2 and SEQ ID NO:12.
 - 28. A recombinant molecule comprising an isolated

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nucleic acid molecule as set forth in Claim 15 operatively linked to a transcription control sequence.

- 29. A recombinant cell comprising a cell having an isolated nucleic acid molecule as set forth in Claim 15, said cell being capable of expressing said nucleic acid molecule.
- 30. The nucleic acid molecule of Claim 15, wherein said nucleic acid molecule comprises a nucleic acid molecule selected from the group consisting of nCtCal₂₅₇, nCtCal₈₅₀, nCtCal₅₅₆, nCtCal₁₅₈₉, nCtCal₁₂₀₉, nCtCal₁₂₁₂, nCtCal₆₆₅, nCtCal₇₅₀, and nCtCal₁₂₁₈.

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- 31. A therapeutic composition for protecting an hematophagous insect infestation, from animal composition comprising a compound selected from the group consisting of an isolated hematophagous insect calreticulin. protein or a mimetope thereof, an isolated nucleic acid hybridizing stringent molecule capable of under hybridization conditions with a flea calreticulin gene, an isolated anti-hematophagous insect calreticulin antibody or a mimetope thereof, a calreticulin inhibitory compound and a mixture thereof, said composition, when administered to an animal, being able to reduce hematophagous insect burden on said animal and in the environment of said animal.
- 32. The composition of Claim 31, wherein said animal is selected from the group consisting of mammals and birds.
- 33. The composition of Claim 31, wherein said animal is selected from the group consisting of cats, dogs, sheep, cows, pigs, horses and goats.

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- A method to protect an animal from hematophagous insect infestation, comprising treating an animal with a therapeutic composition that includes a compound selected from the group consisting of an isolated hematophagous 5 insect calreticulin protein or a mimetope thereof, an isolated nucleic acid molecule capable of hybridizing under stringent hybridization conditions with a flea calreticulin gene, an isolated anti-hematophagous insect calreticulin antibody or a mimetope thereof, a calreticulin inhibitory 10 compound, and a mixture thereof, said composition, when administered to an animal, being able to reduce hematophagous insect burden on said animal and in the environment of said animal.
- 35. The method of Claim 34 or the therapeutic composition of Claim 31, wherein said composition further comprises a component selected from the group consisting of a pharmaceutically acceptable excipient, an adjuvant, a carrier, and a mixture thereof.
 - 36. A method to produce a hematophagous insect calreticulin protein comprising culturing in an effective medium a recombinant cell transformed with a nucleic acid molecule encoding said protein to produce said protein.
 - 37. The method of Claim 36, wherein said nucleic acid molecule comprises a nucleic acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:11 and SEQ ID NO:14.
 - 38. The method of Claim 36, wherein said nucleic acid molecule encodes a protein comprising an amino acid

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sequence selected from the group consisting of SEQ ID NO:2 and SEQ ID NO:12.

- 39. A method to identify a compound capable of inhibiting hematophagous insect calreticulin activity, said method comprising:
- (a) contacting an isolated hematophagous insect calreticulin protein with a putative inhibitory compound under conditions in which, in the absence of said compound, said protein has calreticulin activity; and
- 10 (b) determining if said putative inhibitory compound inhibits said calreticulin activity.
 - 40. A test kit to identify a compound capable of inhibiting hematophagous insect calreticulin activity, said test kit comprising an isolated hematophagous insect calreticulin protein having calreticulin activity and a means for determining the extent of inhibition of said activity in the presence of a putative inhibitory compound.

INTERNATIONAL SEARCH REPORT

International application No. PCT/US96/03133

A. CLASSIFICATION OF SUBJECT MATTER IPC(6): Please See Extra Sheet. US CL: Please See Extra Sheet. According to International Patent Classification (IPC) or to both national classification and IPC B. FIELDS SEARCHED Minimum documentation searched (classification system followed by classification symbols) U.S.: 424/185.1, 275.1; 435/69.1, 252.3, 240.1, 320.1; 514/2, 12; 530/300, 324, 858; 536/23.5 Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched					
Electronic of	data base consulted during the international search (DSIS, MEDLINE, EMBASE erms: flea, calreticulin, antigen				
C. DOC	CUMENTS CONSIDERED TO BE RELEVANT				
Category*	Citation of document, with indication, where	appropriate, of the relevant passages	Relevant to claim No.		
Y	K.K. Murthy et al. Structural h calreticulin gene produce and antigen Ral-1. Nucleic Acids Re page 4933, see entire document.	the Onchocerca volvulus s. 1990, Vol. 18, No. 16,	1, 14-17, 19, 20, 24, 28, 29, 46		
Y	M.J. Smith et al. Multiple zo calreticulin (CRP55, calregulin, libinding ER/SR protein. EMBO Jour pages 3581-3586, especially abs	HACBP), a major calcium rnal. 1989, Vol. 8, Nol. 12,	1, 14-17, 19, 20, 24, 28, 29, 36		
Α	Journal of Cellular Biochemistry issued 1995, abstract no. C: "Characterization of calreticulin in	3-213, Jaworski, D. C.	1-40		
Furthe	er documents are listed in the continuation of Box C	See patent family annex.			
'A' doc	cial categories of cited documents: unent defining the general state of the art which is not considered a of particular relevance	"T" later document published after the inter- date and not in conflict with the applicat principle or theory underlying the inves-	ion but cited to undersiand the ation		
'L' doct	er document published on or after the international filing date ament which may throw doubts on priority claim(s) or which is I to establish the publication date of another citation or other ind reason (as specified)	"X" document of particular relevance; the considered novel or cannot be considered when the document is taken alone "Y" document of particular relevance; the considered to involve an inventive a	ed to involve an inventive step		
men	ament referring to an oral disclosure, use, exhibition or other ns	combined with one or more other such being obvious to a person skilled in the	documents, such combination		
the priority date claimed		*&* document member of the same patent for			
Oate of the a	etual completion of the international search	Date of mailing of the international sear	ch report		
Name and mailing address of the ISA/US Commissioner of Patents and Trademarks Box PCT Washington, D.C. 20231		Authorized officer Durand	Frieder		
acsimile No	. (703) 305-3230	Telephone No. (703) 308-0196			

INTERNATIONAL SEARCH REPORT

International application No. PCT/US96/03133

A. CLASSIFICATION OF SUBJECT MATTER: IPC (6):					
C07K 14/09, 14/435, 16/18; C12N 1/21, 15/09, 15/12, 15/63; A61K 35/64, 38/16, 39/00, 39/35					
A. CLASSIFICATION OF SUBJECT MATTER: US CL :					
424/185.1, 275.1; 435/69.1, 252.3, 240.1, 320.1; 514/2, 12; 530/300, 324, 858; 536/23.5					
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